

DEPARTMENT OF THE AIR FORCE AIR FORCE RESEARCH LABORATORY WRIGHT-PATTERSON AIR FORCE BASE OHIO 45433

5 June 2001

MEMORANDUM FOR US EPA

NCEA (MD-52) RTP, NC 27711

ATTN: ANNIE M. JARABEK

FROM: Elaine Merrill AFRL/HEST

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SUBJECT: Consultative Letter, AFRL-HE-WP-CL-2001-0008, PBPK Model for Perchlorate-Induced Inhibition of Radioiodide Uptake in Humans.

- 1. This letter updates a physiologically-based pharmacokinetic (PBPK) model for predicting the inhibition of thyroid iodide uptake in humans after exposure to perchlorate (ClO₄), as presented in the Consultative Letter, AFRL-HE-WP-CL-2000-0036, Human PBPK Model for Perchlorate Inhibition of Iodide Uptake in the Thyroid. The current model reflects a new model structure and incorporates unpublished data courtesy of Dr. M. Greer (Oregon Health Science University, Portland, OR), Dr. G. Goodman (Intertox, Inc., Seattle, WA), Dr. G. Brabant and Dr. H. Leitolf (Medizinische Hochschule, Hanover, Germany). Nonlinear saturable uptakes are described for the thyroid, skin and stomach compartments. The stomach and thyroid consist of three subcompartments representing the stroma, follicle and colloid in the thyroid, or the capillary bed, stomach wall and contents in the case of the stomach. The skin contains two sub-compartments for capillaries and skin tissue. Compartments for plasma, kidney, liver, fat, richly perfused and slowly perfused tissues were described using passive diffusion. The plasma compartment for perchlorate is composed of plasma and plasma proteins to simulate binding, whereas the plasma compartment for iodide does not include binding.
- 2. The model adequately simulates serum concentrations and cumulative urine after drinking water exposure to perchlorate spanning four orders of magnitude (0.02 to 12.0 mg/kg-day). In addition, the model adequately simulates perchlorate-induced iodide inhibition in the thyroid from doses ranging from 0.007 to 0.5 mg/kg-day. Inhibition data from higher doses were not available.

3. For further information, please contact me by phone: (937) 255-5150 ext. 3195, fax: (937) 255-1474 or e-mail: elaine.merrill@wpafb.af.mil.

ELAINE A. MERRILL
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Attachment:

- 1. PBPK Model for Perchlorate-Induced Inhibition of Iodide Uptake in the Human
- 2. Goodyear, C.: Serum Hormones (TSH, T₃, T₄, fT₄) Statistical Report

1st Ind, AFRL/HEST

5 June 2001

MEMORANDUM FOR US EPA

ATTN: MS. ANNIE JARABEK

This letter report has been coordinated at the branch level and is approved for release.

Richard R. Stotts, DVM, Ph.D.

Branch Chief

Operational Toxicology Branch Human Effectiveness Directorate

PBPK Model for Perchlorate-Induced Inhibition of Radioiodide Uptake in Humans

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5 June 2001

INTRODUCTION

Commercially the salt, ammonium perchlorate, has been produced in the U.S. since the 1940s (Fisher *et al.*, 2000). It is used primarily for its strong oxidizing properties in explosive and pyrotechnic mixtures and as an ingredient in rocket fuel. In water, perchlorate quickly dissociates and is highly mobile and stable. As a result, perchlorate contamination has been detected in several water sources in the U.S. (Urbansky and Schock, 1999). There's concern that chronic exposure to low levels of perchlorate in drinking water could induce iodide deficiencies and subsequent thyroid disorders.

The perchlorate anion (ClO₄) is very similar in ionic size, shape and charge to iodide (Γ). These shared properties allow perchlorate to interfere with the first stage of thyroid hormone genesis by competitively inhibiting the active transfer of iodide into the thyroid by the sodium-iodide symporter (NIS). The NIS is a protein that resides in the basolateral membrane of thyroid epithelial cells (Spitzweg *et al.*, 2000). NIS simultaneously transports both Na⁺ and Γ ions from extracellular fluid (plasma) into the thyroid epithelial cell. This process is an example of secondary active transport. Energy is provided by the electrochemical gradient of sodium across the cell membrane; the low intracellular concentration of sodium is maintained by sodium-potassium pumps (Ajjan *et al.*, 1998).

The presence of NIS is an indicator of active uptake for iodide and likewise, perchlorate. The NIS is highly expressed in thyroid epithelial cells. Lower levels of expression have also been detected in the mammary gland, salivary gland, skin, stomach and colon. Perchlorate inhibits iodide uptake in these tissues as well. However, only the thyroid has been found to organify iodide to form thyroid hormones (Ajjan *et al.*, 1998; Spitzweg *et al.*, 1998). The perchlorate-induced decline in iodide trapping in all tissues with NIS is reversible upon elimination of perchlorate.

Thyroid hormones homeostasis occurs through a complex feedback mechanism. Thyroid-stimulating hormone (TSH) is the most important regulator of NIS gene and protein expression; up-regulation of NIS stimulates iodide transport for increased hormone production. A drop in thyroid hormones signals the pituitary to produce more TSH, which in turn stimulates NIS expression. In rats, an increase in TSH and decrease in free thyroxine (fT₄) occurs quickly after perchlorate exposure (Wyngaarden *et al.*, 1952). In humans, however, the conservation of thyroid hormones is more efficient. Studies have shown little or no change in human TSH levels after two weeks of exposure perchlorate in drinking water (Greer *et al.*, 2000; Brabant *et al.*, 1992). Therefore, the extent of chronic low level perchlorate exposure required to cause significant hormone deficiencies is not yet known.

Adequate iodide uptake for thyroid hormone production is necessary for proper brain development during the first two years of life (Laurberg *et al.*, 2000). Iodide deficiency during the fetal and neonatal period could result in abnormal physical and mental development (Porterfield, 1994). The effects of iodide deficiencies to adult populations are less dramatic.

However, in elderly adults, especially elderly women, a high occurrence of hyperthyroidism is noted in areas of mild to moderate iodine deficiency. Hyperthyroidism is easily overlooked in the elderly and, if left untreated, may lead to cardiac arrhythmias, impaired cardiovascular reserves, osteoporosis and other abnormalities (Laurberg *et al.*, 2000). With an aging population, this may become a greater public health concern.

Currently, perchlorate is not regulated in the U.S. under the Federal Safe Drinking Water Act. Occupational health studies at two perchlorate production facilities in the U.S., in which perchlorate exposure was quantified through urinary perchlorate measurements, found no elevation in TSH, drop in fT₄ or other adverse effects (Gibbs *et al.*, 1998). Epidemiological studies on neonatal screening data from California and Nevada health departments showed no increase in incidence of congenital hypothyroidism or decrease in neonatal T₄ associated with perchlorate in drinking water up to 15 μ g/L (Lamm and Doemland, 1999; Li *et al.*, 2000). Serum samples were collected from school-age children and newborns in three cities in Chile with reported perchlorate in drinking water (<4, 5 to 7 and 100 to120 μ g/L). Samples revealed significantly higher free T₄ but normal TSH levels in the two cities with higher perchlorate concentrations (Crump *et al.*, 2000). Dietary iodide in each of the above populations was within normal ranges.

The proposed model describes the kinetics and distribution of perchlorate in the average adult human after exposure to the perchlorate. It also describes iodide kinetics and competitive inhibition of iodide uptake by perchlorate. Perchlorate's transport mechanisms can be modeled similarly to that of iodide, as it binds to the NIS and competitively inhibits iodide uptake (Anbar et al., 1959; Brown-Grant and Pethes, 1959). The kinetics of these anions differ mainly in that iodide is organified in the thyroid (thyroid hormone production) whereas perchlorate is thought to be unreactive and eventually diffuses from the thyroid into systemic circulation. Although perchlorate is quickly eliminated unchanged in the urine (Wolff, 1998), the impact of chronic displacement of iodide from prolonged exposure to perchlorate-contaminated drinking water is the focus of this and ongoing modeling efforts.

The objective of this effort is to simulate serum perchlorate and iodide levels and the subsequent inhibition of iodide uptake into the thyroid. The model does not include effects on thyroid hormone production and homeostasis at this point.

METHODS

The human data used in model development were obtained from Hays and Solomon (1965) and both published and unpublished data from a recent human study involving drinking water exposure to perchlorate and measured inhibition of radioiodide uptake in the thyroid (Greer *et al.*, 2000).

Data supporting model validation were obtained from another recent but unpublished drinking water study conducted by Drs. Holger Leitolf and Georg Brabant of the Medizinische Hochschule, Hanover, Germany. In addition, urinary perchlorate clearance data by Eichler

(1929), Kamm and Drescher (1973) and Durand (1938) were also used to validate model predictions.

Human Iodide Kinetic Data (Hays and Solomon, 1965)

A comprehensive human kinetic study on early iodide distribution was reported in 1965 by Hays and Solomon. The authors studied the effect of gastrointestinal cycling on iodide kinetics in nine healthy males after an intravenous (*iv*) dose of 3.44 x 10⁻³ ng ¹³¹ Γ /kg bodyweight. Frequent measurements of radioiodide uptake in the thyroid, gastric secretions, plasma and cumulative urine samples were taken during the three hours following injection. Gastric secretions were collected using a nasogastric tube with constant suction while the subjects remained in a resting position (only standing to urinate). Saliva was not collected separately and therefore pooled, to some extent, with gastric juices. To account for the removal of gastric iodide from circulation and to determine its impact on free iodide distribution, the authors ran a control session on the same subjects without aspirating gastric secretions. Aspirated gastric secretions accounted for 23% of the ¹³¹ Γ administered.

Perchlorate Kinetics and Inhibition of Thyroid Iodide Uptake (Greer et al., 2000)

Perchlorate Data

Greer *et al.* (2000) recently studied effects on humans of repeated low level exposure to perchlorate. Subjects received 0.5, 0.1, 0.02 or 0.007 mg/kg-day perchlorate in drinking water over a two week period. Each dose group consisted of eight healthy volunteers (four males and four females), with no signs or symptoms of thyroid disorders. The daily dose was dissolved in 400 mL water and divided into four 100 mL servings, which were ingested at approximately 0800, 1200, 1600 and 2000 hours.

Baseline serum and urine samples were collected before the first perchlorate treatment. During perchlorate exposure, serum samples were collected at the following approximate times: day 1 at 1200 and 1600, day 2 at 0800, 1200 and 1700, day 3 at 0900, day 4 at 0800 and 1200, day 8 between 0800 and 0900 and day 14 at 0800 and 1700. Serum samples were also collected on post-exposure days 1, 2, 3 and 14. Twenty-four hour urine collections were taken on exposure days 1, 2, 14 and post-exposure days 1 through 3. Serum and 24 hour urine samples from the study were provided compliments of Dr. Monte Greer of Oregon Health Science University (OHSU), Portland, OR, and Dr. Gay Goodman of Intertox, Seattle, WA; samples were analyzed for perchlorate at the Operational Toxicology Branch, Human Effectiveness Directorate at the Air Force Research Laboratory (AFRL/HEST), Wright Patterson Air Force Base (WPAFB), OH, using the analytical methods described below. A detailed description of the study method is provided in Greer *et al.* (2000).

Iodide Inhibition Data

Eight and 24 hour thyroid ¹²³Γ uptakes (radioiodine uptake or RAIU) were measured one to two days prior to perchlorate treatment (baseline), on days 2 and 14 of perchlorate exposure and 14 days after perchlorate. A gelatin capsule containing 100 μCi of ¹²³Γ was administered orally at 0800, before drinking the first perchlorate solution for that day. Thyroid scans were then taken 8 and 24 hours later.

Thyroid Hormone Data

The serum samples were also analyzed for TSH, T₄, T₃ and fT₄ at OHSU. These hormone data were not used in the PBPK model described below; however, statistical analysis of the data is described in Attachment 2.

Supporting Kinetic Studies

Both urine and serum perchlorate concentrations were provided from a recent unpublished study by Drs. Brabant and Leitolf of Medizinische Hochschule, Hanover, Germany. In their study, 7 healthy males ingested 12.0 mg/kg-day perchlorate dissolved in 1 liter of drinking water everyday for 2 weeks. The daily perchlorate dose was divided equally in three portions and ingested three times per day (approximately between 0600 and 0800, 1100 and 1300, and 1800 and 2000 hours). Blood specimens were collected on days 1, 7 and 14 of perchlorate treatment and on the two mornings after perchlorate administration was discontinued. Samples were analyzed for perchlorate at AFRL/HEST.

Three published studies reported cumulative urine concentrations collected from healthy males after receiving a high oral dose of perchlorate (Durand, 1938; Kamm and Drescher, 1973; Eichler, 1929). Oral doses administered in these studies were 784 mg NaClO₄ (635 mg ClO₄) (Durand, 1938); 1000 mg NaClO₄ (765 mg ClO₄) (Kamm and Drescher, 1973) and 2000 mg KClO₄ (1400 mg ClO₄) (Eichler, 1929). The studies did not report serum perchlorate levels.

Stanbury and Wyngaarden (1952) measured radioiodide uptake in a patient with Grave's disease. The patient received a tracer dose of ¹³¹ as a control before perchlorate dosing and again one hour after administration of 100 mg KClO₄. Thyroid scans of radioiodide uptake were performed both after the control and perchlorate sessions to determine the level of inhibition.

Analytical Methods

Serum samples were analyzed for perchlorate by ion chromatography on a Dionex DX-500 ion chromatography system with a GP-40 gradient pump, CD-20 conductivity detector, a LC-20 chromatography enclosure and an AS40 automated sampler. The injection volume was 200 μ L. Anion separation was obtained on a Dionex Ion Pac AS-11, 2.0 x 250-mm separation column

with an AG-11 2.0 x 50 mm guard column and an ATC-1 anion trap column. The mobile phase consisted of 80 mM NaOH. The mobile phase flow rate was set at 0.25 mL/min. Background suppression was achieved by using an Anion Self-Regenerating Suppressor (ASRS)-ULTRA suppressor, with external water flowing at 10 mL/min.

For sample preparation, $50~\mu L$ of serum was precipitated with $200~\mu L$ of cold 100% ethanol. Samples were then centrifuged at 14,000~rpm for 30~minutes, using an Eppendorf microcentrifuge. The supernatant was removed and evaporated to dryness under the flow of nitrogen gas. Samples were then reconstituted in 1~mL of $18~\text{M}\Omega/\text{cm}$ water. The reconstituted samples were filtered through a Millipore Millex HV-13 0.45~micron syringe filter and placed in 2~mL sample vials for analysis. The samples required no further dilution, making the final dilution after preparation 1:20. To check the performance of the instrument, a duplicate sample, a perchlorate spiked sample and control standards were evaluated after every ten serum samples.

Ion chromatography of urine was performed on a Dionex DX-500 microbore ion chromatograph system with a GP-40 gradient pump, CD-20 conductivity detector, an LC-20 chromatography enclosure and an AS40 automated sampler. The injection volume was 200 μ L. Anion separation was obtained on a Dionex IonPac AS-11 2.0 x 250-mm separation column with an AG-11 2.0 x 50 mm guard column and an ATC-1 anion trap column. The mobile phase varied from 60 to 120 mM NaOH, depending on the sample. The mobile phase flow rate was set at 0.25 mL/min. Background suppression was achieved by using an Anion Self-Regenerating Suppressor (ASRS)-ULTRA suppressor, with external water flowing at 10 mL/min.

For the sample preparation of urine, $500~\mu L$ of urine was filtered through a Millipore Millex HV13 0.45 micron syringe filter. The sample was then placed in a Millipore Ultrafree 0.5 centrifuge filter and was centrifuged in an Eppendorf microcentrifuge at 14,000 rpm for 30 minutes. The filtered sample was then removed and diluted 1:500 in 2 mL sample vials. The samples in which no perchlorate was detected were prepared a second time, using the same process as described above, but diluted only 1:50. If there was still no perchlorate detected in the samples, then they again were prepared with the same method as above and diluted 1:25. A 1:25 dilution is the lowest dilution that can be used without producing a significant increase in baseline noise and a subsequent decrease in detection capabilities.

Model Structure

The human model for perchlorate inhibition of iodide uptake was developed concurrently with the male rat model (Merrill, 2001). Much of the early development was based upon generalizations drawn from in-house work on perchlorate (Fisher *et al.*, 2000) and the work of Hays and Wegner (1965) describing iodide kinetics. Nearly identical model structures were used to describe the kinetics of both iodide and perchlorate, as they share similar ionic size and an affinity for NIS (Figure 1). Tissues, which were reported to have tissue:plasma concentration ratios greater than one and that express NIS were described as compartments of nonlinear saturable uptake. In the model, these tissues include the thyroid, skin and gastric mucosa (Wolff, 1998; Chow *et al.*, 1969; Kotani *et al.*, 1998; Anbar *et al.*, 1959 and Perlman *et al.*, 1941).

Although other tissues have been known to sequester iodide and similar anions (e.g., salivary glands, choroid plexus, ovaries, mammary glands, placenta) (Brown-Grant, 1961; Honour *et al.*, 1952; Spitzweg *et al.*, 1998), the iodide and perchlorate pools of these tissues are expected to be too small to significantly affect plasma levels or were not important in the non-pregnant human. These tissues were therefore lumped with slowly and richly perfused tissues.

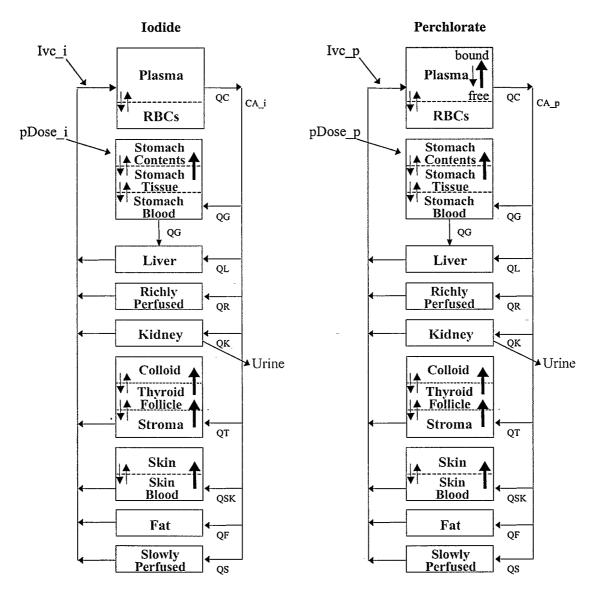


Figure 1. Schematic of PBPK model for perchlorate and iodide distribution. Bold arrows indicate active uptake at NIS sites (with exception of plasma binding). Small arrows indicate passive diffusion.

In addition to the thyroid, stomach, skin, slowly and richly perfused compartments, the model also includes separate compartments for plasma, kidney, liver and fat. The kidney was required

to describe urinary clearance of the anions. A liver compartment was maintained separately, as it would be required in future development of the model to include hormone homeostasis. The liver is the primary site of extrathyroidal deiodination. Fat does not absorb either anion well. However, wide variations in percent body fat, between sexes and individuals, in human populations could alter perchlorate/iodide kinetics. Therefore, for the purpose of using the model for species extrapolations, fat is included as an exclusionary compartment.

The stomach and thyroid consist of three sub-compartments representing the stroma, the follicle and the colloid in the thyroid, or the capillary bed, stomach wall and contents in the case of the stomach. The skin contains two sub-compartments representing the capillary bed and the skin tissue. Active uptake into the thyroid colloid, stomach contents and skin were described using Michealis-Menten kinetics for nonlinear processes (Figure 1, bold arrows). Permeability area cross products (PAs) and partition coefficients (Ps) were used to describe the first order movement of the anions between the capillary bed, tissue and inner compartments (Figure 1, small arrows), which results from the inherent electrochemical gradient within the tissues. Passive diffusion through the kidney, liver and fat compartments were described with partitions and blood flows for these tissues. Plasma binding of perchlorate was described with Michealis-Menten terms for the association of the ClO₄⁻ anions to plasma binding sites and a first order clearance rate for the dissociation. First order clearance rates from the kidney were also used to describe urinary clearance of the anions.

There are few studies that measured perchlorate or iodide in skin. Yu et al. (2000) reported ³⁶ClO₄ in male rat skin at higher concentrations than in plasma for several hours after *iv* administration. Zeghal et al. (1995) examined the effects of perchlorate on the iodine composition of rat skin. Although Zeghal and his colleagues did not measure skin perchlorate levels directly, they reported a significant reduction in skin iodide in young and adult rats after its administration, suggesting competitive inhibition. Therefore, active uptake was also used to describe the skin compartment for perchlorate.

Iodide has also been reported in the skin at higher concentrations than in plasma after *iv* dosing, but the results have not been consistent. Brown-Grant and Pethes (1959) observed skin:serum ratios as high as 2.0 in adult male rats and 5.0 in young pups. The skin:serum ratio was higher in the adult male than female rats. They did not find skin:serum ratios higher than one in guinea pigs or mice, however. *In vivo* studies with radioiodide in rats resulted in skin:serum iodide ratios near one (Yu *et al.*, 2000), but not at levels as high as those reported by Brown-Grant and Pethes.

This behavior in skin has been reported for another similar anion. Hays and Green (1973) performed dialysis studies on intact human tissues with pertechnetate. They found skin had a relatively slow uptake of pertechnetate peaking at 18 hours and, in fact, more retention after leaching dialysis than brain, muscle and serum.

Clinical scans of radioiodide tracers have not revealed elevated iodide levels in human skin. This does not necessarily imply that iodide is not actively sequestered to some extent in human skin. It is possible that skin radioactivity was diffused enough to be indistinguishable from

background. The fact that NIS expression has been reported in rat skin (Kotani *et al.*, 1998) and the distribution of NIS has been found to be similar across species (Wolff, 1998) suggests that the skin, due to its size, may represent an important pool for slow iodide turnover in humans.

The blood compartment differs between the perchlorate and iodide models. The perchlorate blood compartment is composed of plasma and plasma proteins to simulate binding. Free iodide does not appear to bind to plasma proteins in the same manner as ClO₄ and therefore a single compartment for plasma is used in the iodide model.

Physiological Parameters

Human tissue volumes and blood flows were obtained from the literature (Table 1). Considerable variability was reported for some parameters. For example, blood flow to the gastrointestinal (GI) tract can increase tenfold in response to enhanced functional activity (secretion and digestion) (Granger *et al.*, 1985). Blood flows used in the model represent estimates of resting values. Human data on the volume of the stomach capillary bed (VGBc) were not found in the published literature. Therefore, a value derived from rat stomach data (Altman and Dittmer, 1971a) was applied as VGBc in the human.

Thyroid volume was obtained from ultrasound measurements on 57 healthy volunteers with no thyroid disorders (37 to 74 years of age), in a study conducted by Yokoyama $et\ al.$ (1986). The mean thyroid volume was 13.4 ± 4.1 mL and mean thyroid volume to bodyweight ratio was 0.251 ± 0.074 mL/kg (mean \pm standard deviation (s.d.)), approximately 0.03% of bodyweight (BW). They found a positive correlation between thyroid volume and both bodyweight and age, with weight having the most pronounced influence. The percent of total thyroid volume attributed to the thyroid follicular epithelium, colloid and stroma were estimated from histometric measurements of patients at necropsy by Brown $et\ al.$ (1986). Their findings on the histological features of thyroids of men and women showed overlapping distributions without evidence of a significant difference between sexes.

A significant sex difference in total fat mass is reported in humans, with women having approximately 10% more fat than men (Brown *et al.*, 1997). Therefore, a gender-specific value was used for this parameter.

Table 1. Human physiological parameters

Physiological Parameters	Human	Source		
Tissue Volumes				
Bodyweight BW (Kg)	~70.0	Subject-specific		
Slowly Perfused VSc (%BW)	65.1	Brown et al., 1997		
Richly Perfused VRc (%BW)	12.4	Brown et al., 1997		
Fat VFc (%BW)	♂ 21.0	Brown et al., 1997		
	Q 32.7			
Kidney VKc (%BW)	0.44	Brown et al., 1997		
Liver VLc (%BW)	2.6	Brown et al., 1997		
Stomach Tissue VGc (%BW)	1.7	Brown et al., 1997		
Gastric Juice VGJc (%BW)	0.071	Licht and Deen, 1988		
Stomach Blood VGBc (%VG)	4.1	Altman & Dittmer, 1971a		
Skin Tissue VSkc (%BW)	3.7	Brown et al., 1997		
Skin Blood VSkBc (%VSk)	8.0	Brown et al., 1997		
Thyroid VTtotc (%BW)	0.03	Yokoyama et al., 1986		
Thyroid Follicle VTc (%VTtot)	57.3	Brown et al., 1986		
Thyroid Lumen VDTc (%VTtot)	15.0	Brown et al., 1986		
Thyroid Blood VTBc (%VTtot)	27.6	Brown et al., 1986		
Plasma Vplasc (%BW)	4.4	Marieb, 1992; Altman & Dittmer, 1971b		
Red Blood Cells VRBCc (%BW)	3.5	Marieb, 1992; Altman & Dittmer, 1971b		
Adjusted Slowly Perfused VS (L)	28.0	Calculated from model		
Adjusted Richly Perfused VR (L)	5.34	Calculated from model		
Blood Flows				
Cardiac Output QCc (L/hour-kg)	16.5	Brown et al., 1997; Hanwell & Linzell, 1973		
Fat QFc (%QC)	5.2	Brown et al., 1997		
Kidney QKc (%QC)	17.5	Brown et al., 1997		
Liver QLc (%QC)	22.0	Brown et al., 1997		
Stomach QGc (%QC)	1.0	Leggett & Williams, 1995; Malik et al., 1976		
Thyroid QTc (%QC)	1.6	Brown et al., 1997		
Adjusted Slowly Perfused QS (%QC)	13.0	Calculated, using 24% QC as flow to all		
		slowly perfused tissues (Brown et al., 1997)		
Adjusted Richly Perfused QR (%QC)	33.0	Calculated, using 76% QC as flow to all		
	<u> </u>	richly perfused tissues (Brown et al., 1997)		

Partitioning Coefficients

Partition coefficients for iodide and perchlorate were estimated from *in vivo* studies (Table 2). Tissue and serum measurements from steady state conditions were not available. Therefore, they were estimated from the clearance portion of the data collected after acute dosing. Halmi *et al.* (1956) measured organ to serum concentration ratios for radioiodide in rats approximately 1, 4 and 24 hours after an *iv* dose of the tracer iodide. The average liver:serum and muscle:serum iodide ratios at approximately 4 hour after an injection of ¹³¹ Γ (0.40 and 0.21, respectively) were used to represent human rapidly and slowly perfused partitioning coefficients for iodide.

Perlman et al. (1941) reported similar iodine ratios in rabbit tissues five hours after subcutaneous dosing with NaCl and a tracer amount of iodide (0.44 for liver/blood and 0.19 for muscle/blood). These liver:blood and muscle:blood, ratios reported by Perlman remained relatively constant for up to 96 hours.

Perchlorate partition coefficients for rapidly (0.56) and slowly perfused (0.31) tissues were derived from in-house rat studies, 24 hours after a single *iv* dose of 3.3 mg ³⁶ClO₄-/kg (Yu *et al.*, 2000). Anbar *et al.* (1959) reported similar liver:blood and muscle:blood ratios of 0.38 and 0.12, respectively, in rabbits 12 hours after an intraperiteneal dose of 100 mg KClO₄.

A value of 0.05 was reported for the partitioning of perchlorate into the fat of a hen (Pena et al., 1976). This value was also used to represent iodide partitioning into fat in our model, since other tissue:serum concentration ratios from Pena's hen study were consistent with ratios found in both the rat (Yu et al., 2000) and rabbit (Anbar et al., 1959).

For compartments with nonlinear uptake of the anions, effective partition coefficients were used, representing either approximate tissue:serum concentration ratios or electrical potential gradients. The effective partition coefficients for iodide in the stomach compartments were also derived from in-house rat experiments. Stomach wall:serum and gastric juice:stomach wall perchlorate ratios of 1.8 and 2.3, respectively, were derived 24 hours after the *iv* dose. Estimated stomach iodide partitions, derived from in-house ¹²⁵I kinetic studies, were 1.0 for the stomach tissue:serum and 3.5 for the gastric juice:stomach ratios(Yu, 2000).

Skin measurements from the in-house ³⁶ClO₄ kinetic study were highly variable and suggested an effective partitioning greater than one. Based on the 24 hour timepoint, an effective partition coefficient of 1.15 was derived (Yu, 2000). The iodide value used (0.7) was derived from Perlman *et al.* (1941). They reported skin:plasma iodide ratios from 0.6 to 0.7 in rabbits, 6 to 96 hours after a subcutaneous dose of tracer radioiodide. This is in agreement with in-house ¹²⁵I studies, which suggest a partition coefficient less than 1.0 (Yu *et al.*, 2000).

Chow and Woodbury (1970) measured electrochemical potentials within the thyroid stroma, follicular membrane and colloid at three different doses of ClO₄. Their measured difference in electrical potential between the thyroid stroma and follicle can be interpreted as an effective partition coefficient for charged moieties, such as ClO₄ and Γ, hindering the entry of negatively charged ions into the follicle. The equal and opposite potential from the follicle to the colloid enhances passage of negatively charged species into the colloid and indicates an effective partition coefficient of greater than one.

From Chow and Woodbury (1970), the potential difference for the stroma: follicle interface ranges from -58 to -51 mV, from which an effective partitioning between 0.114 and 0.149 is calculated (Clewell, 2001a; Merrill, 2001). Similarly, the follicle:lumen potential ranges from +50 to +58 mV, resulting in an effective partition between 6.48 and 8.74. These values were also used to describe the uptake of iodide based on the fact that iodide and perchlorate have the same ionic charge (-1) and similar ionic radii and therefore react similarly to the electrochemical gradient.

Table 2. Chemical specific parameters for human model

Partition Coefficients (unitless) Ps	Iodide	Perchlorate	Source	
Slowly Perfused / Plasma PS_	0.21	0.31	Halmi et al., 1956;Yu et al., 2000	
Richly Perfused / Plasma PR_	0.40	0.56	Halmi et al., 1956;Yu et al., 2000	
Fat/ Plasma PF_	0.05	0.05	Pena et al., 1976	
Kidney/ Plasma PK	1.09	0.99	Perlman et al., 1941; Yu et al., 2000	
Liver/Plasma PL	0.44	0.56	Perlman et al., 1941; Yu et al., 2000	
Gastric Tissue/Gastric Blood PG	0.50	1.80	Yu et al., 2000;Yu., 2000	
Gastric Juice/Gastric Tissue PGJ	3.50	2.30	Yu et al., 2000; Yu, 2000	
Skin Tissue/Skin Blood PSk	0.70	1.15	Perlman et al., 1941; Yu, 2000	
Thyroid Tissue/Thyroid Blood PT	0.15	0.13	Chow & Woodbury (1970)	
Thyroid Lumen/Thyroid Tissue PDT	7.00	7.00	Chow & Woodbury (1970)	
Red Blood Cells/Plasma	1.00	0.80	Rall et al., 1950; Yu et al., 2000	
Max Capacity, Vmaxc's (ng/hr-kg)				
Thyroid Colloid Vmaxc DT	1.0E+8	2.5E+5	Fitted	
Thyroid Follicle Vmaxc T	~1.5E+5	5.0E+4	Fitted	
Skin Vmaxc_S	7.0E+5	1.0E+6	Fitted	
Gut Vmaxc_G	9.0E+5	1.0E+5	Fitted	
Plasma binding Vmaxc Bp		5.0E+2	Fitted	
Affinity Constants, Km's (ng/L)				
Thyroid Lumen Km_DT	1.0E+9	1.0E+8	Golstein et al., 1992	
Thyroid Km_T	4.0E+6	1.8E+5	Gluzman & Niepomniszcze, 1983; Wolff, 1998	
Skin Km_S	4.0E+6	2.0E+5	Gluzman & Niepomniszcze, 1983; Wolff, 1998	
Gut Km_G	4.0E+6	2.0E+5	Gluzman & Niepomniszcze, 1983; Wolff, 1998	
Plasma binding Km_B		1.8E+4	Fitted	
Permeability Area Cross Products, P.	As (L/hr-l	κg)		
Gastric Blood to Gastric Tissue PAGc	0.2	0.6	Fitted	
Gastric Tissue to Gastric Juice PAGJc	2.0	0.8	Fitted	
Skin Blood to Skin Tissue PASkc_	0.06	1.0	Fitted	
Plasma to Red Blood Cells PARBCc_	1.0	1.0	Fitted	
Follicle to Thyroid blood PATc_	1.0E-4	1.0E-4	Fitted	
Lumen to Thyroid Follicle PADTc_	1.0E-4	0.01	Fitted	
Clearance Values, Cl's (L/hr-kg)				
Urinary excretion CLUc_	0.1	0.126	Fitted	
Plasma unbinding Clunbc		0.025	Fitted	

Note: All parameters listed are notated in the model by either an *i* (for iodide) or *p* (for perchlorate) following the parameter name (e.g., PR_i , PR_i

Affinity Constants and Maximum Velocities

Gluzman and Niepomniszcze (1983) derived a mean Michaelis-Menten affinity constant (Km) of 4.0×10^6 ng/L for iodide from thyroid slices of 5 normal individuals. The thyroid slices were incubated with several medium iodide concentrations. The authors noted little variation between normal and pathological human thyroid specimens, or between thyroid specimens of different

species. Therefore, a Km value of 4.0×10^6 ng/L was assumed to describe the uptake of iodide in compartments involving active uptake by NIS (thyroid and gastric juices).

Wolff (1998) noted that iodide's Km for NIS is similar in different tissues. Wolff also noted that the Km for perchlorate and other similar monovalent anions decreased with the anions' ability to inhibit iodide uptake. Several studies suggest perchlorate is a more potent inhibitor than iodide. In the rat thyroid, Wyngaarden *et al.* (1952) have shown that perchlorate was a more powerful inhibitor of the iodide trap than thiocyanate. Halmi and Stuelke (1959) showed that perchlorate was ten times as effective as iodide in depressing tissue to blood ratios in the rat thyroid and gut. Similarly, Harden *et al.* (1968) compared human saliva to plasma radioiodide concentration ratios after equimolar doses of perchlorate and iodide. The saliva/plasma iodide ratios, during resting conditions, were approximately seven times lower after a molar equivalent dose of perchlorate vs. iodide. Lazarus *et al.* (1974) also demonstrated that perchlorate was taken up to greater extent in mice salivary glands than iodide. Based on this information, a Km between 1.8 x 10⁵ and 2.0 x 10⁵ ng/L, approximately 10 times lower than that of iodide, was estimated to represent perchlorate's higher affinity for NIS.

The apical follicular membrane also exhibits a selective iodide uptake mechanism. Golstein *et al.* (1992) found the Km for iodide transport from the bovine thyroid follicle into the colloid (Km_DTp) to be approximately 4.0×10^9 ng/L. This iodide channel also appears to be very sensitive to perchlorate inhibition. The ability of perchlorate to inhibit iodide uptake at the apical follicular membrane suggests that the Km of perchlorate at the apical follicular membrane (Km_Dtp) is also lower than that of iodide. Model simulations of thyroid inhibition supported a value of 1.0×10^8 ng/L, approximately ten times less than that of iodide for this channel also.

Whereas the Km is similar across tissues containing NIS, the maximum velocity term (Vmax) does vary between tissues and species (Wolff, 1998), being lower in humans than other species (Gluzman and Niepomniszcze, 1983; Wolff and Maurey, 1961) when expressed per gram of tissue. Maximum velocities or capacities (Vmaxc) were not found in the literature and were therefore estimated for a given compartment by fitting the simulation to the data at varying doses (Table 2).

TSH increases the total amount of NIS in a membrane, thereby increasing the thyroid Vmax. Gluzman and Niepomniszcze (1983) measured elevated Vmax(s) in thyroid specimens from subjects with Grave's disease, toxic adenoma and dishormonogenetic goiter. In specimens from non-toxic nodular goiter, Hashimoto's thyroid or extranodular tissue from toxic adenoma, maximum capacities were decreased. They found no significant difference between normal human glands and warm nodules from nodular goiters. In addition to hyper- or hypo-stimulation of trapping activity (thyroid disorders), the intrathyroidal iodide pool and the magnitude of iodide efflux are also responsible for variations in Vmax (Bagchi and Fawcett, 1973). Currently the iodide model does not account for TSH stimulation and endogenous iodide pools.

TSH regulation is reported to be unique to the thyroid NIS and does not regulate uptake in other tissues with NIS (Cavalieri, 1997). Spitzweg *et al.* (2000) suggest the up-regulation of NIS protein expression in mammary glands of lactating rats may be regulated by prolactin and/or

oxytocin (Spitzweg *et al.*, 2000), inferring that other agents may stimulate NIS expression and function. The effect of TSH on iodide transport may also be physiologically responsive to other hormones and growth factors. Most growth factors are reported to decrease the ability of TSH to stimulate iodide accumulation, with the exception of insulin, which has an opposite effect (Carrasco, 1993).

Fitting Model Parameters to Experimental Data

Parameter terminology used in this model is summarized in Table 2. Simultaneous differential equations, which simulate radioiodide and perchlorate distribution in the proposed mathematical model, were written and solved using ACSLTM (Advanced Continuous Simulation Language) software (AEgis Technologies, Austin, TX).

Permeability Area Cross Products and Clearance Values

Diffusion limited uptake in tissues requiring subcompartments (e.g., the stomach, thyroid and skin) were described using permeability area cross products (PA) (L/hour-kg) and partition coefficients (P). The PA values in this model were fitted by setting the partition coefficients to the literature values in Table 2. The equations below illustrate the use of PAs:

$$RAXB _ y = QX \times (CA _ y - CVXB _ y) + PAX _ y \times \begin{pmatrix} CX _ y / \\ PX _ y - CVXB _ y \end{pmatrix}$$

$$AXB _ y = \int_{0}^{t} (RAXB _ y) dt$$

$$CVXB _ y = AXB _ y / VXB$$

$$RAX _ y = PAX _ y \times \begin{pmatrix} CVXB _ y - CX _ y / \\ PX _ y \end{pmatrix}$$

$$AX _ y = \int_{0}^{t} (RAX _ y) dt$$

$$CX = AX / VX$$

where:

Net rate of uptake of yth anion (I or ClO₄) into Xth tissue's capillary bed RAXB y =(ng/hour) Amount of yth anion in Xth tissue's capillary bed (ng) AXB y =Blood flow through Xth tissue (L/hour) QX Arterial blood concentration of yth anion (ng/L) CA y = -Net rate of uptake of yth anion into Xth tissue (ng/hour) Amount of yth anion in Xth tissue (ng) $RAX_y =$ AX y Venous blood concentration of yth anion (ng/L) CVXB y =Tissue concentration of yth anion (ng/L) CX y

Fitted clearance values were used to describe first order urinary excretion rates and reversible plasma binding to serum, as shown below:

$$RAX_y = CX_y \times ClX_y$$

where: RAX = rate of clearance of y^{th} anion from X^{th} compartment (ng/hour) $CX_y = \text{concentration of } y^{\text{th}} \text{ anion within } X^{\text{th}} \text{ compartment (ng/L)}$ CIX y = clearance value of y^{th} anion into X^{th} compartment (L/hour)

Saturable Processes

The basic equation used to simulate active uptake of iodide and perchlorate alone (without accounting for inhibition) in tissues with NIS activity is:

$$\frac{dAX - y}{dt} = \frac{V \max_{X} Xy \times CVX - y}{Km_{X}y + CVX_{y}}$$

where:

 $AX_y = Amount of y^{th} anion in X^{th} tissue (ng)$

t = Time (hour)

Vmax_Xy = Maximum uptake of y^{th} anion at X^{th} tissue's symporter (ng/hour) Km_Xy = Michaelis-Menten (M-M) affinity constant for y^{th} anion in X^{th}

compartment (ng/L)

= Concentration of yth anion in capillary blood of Xth compartment (ng/L)

Accounting for inhibition of active uptake of either iodide or perchlorate in the presence of the competing anion is expressed as:

$$\frac{dAX - y}{dt} = \frac{V \max_{Xy \times CVX} y}{Km_Xy \times \left(1 + \frac{CVX}{x} - \frac{z}{Km_Xz}\right) + CVX_y}$$

where:

Km Xz = M-M affinity constant for z^{th} anion (competitive inhibitor) in X^{th} compartment (ng/L)

 $CVX y = Concentration of z^{th}$ anion (competitive inhibitor) in venous capillary blood of Xth compartment (ng/L)

Example equations for simulating the transport of iodide (i) through the thyroid (T) are provided below. The equations include blood flow through the thyroid capillary bed (stroma) as well as active uptake and inhibition by perchlorate in both the follicle and colloid. The transport of the anions through other tissues of active uptake is similar.

• Rate of change in thyroid blood (RATB i)

$$RATB_i = QT \times (CA_i - CVTB_i) + PAT_i \times \begin{pmatrix} CT_i / PT_i - CVTB_i \end{pmatrix} - RupT_i$$

• Rate of change in follicle (RAT i)

$$RAT_i = RupT_i + PAT_i \times \begin{pmatrix} CVTB_i - CT_i \\ PT_i \end{pmatrix} - RupDT_i + PADT_i \times \begin{pmatrix} CDT_i \\ PDT_i - CT_i \end{pmatrix}$$

• Rate of change in Colloid (RADT i)

$$RADT_i = RupDT_i + PADT_i \times \begin{pmatrix} CT_i - CDT_i \\ PDT_i \end{pmatrix}$$

• Rate of active uptake in the follicle $(RupT_i)$ with inhibition by perchlorate (p)

$$RupT _i = \frac{V \max_{Ti} \times CVTB _i}{Km_{Ti} \times \left(1 + \frac{CVTB}{p} - \frac{p}{Km_{Tp}}\right) + CVT_i}$$

Rate of active uptake into the colloid (RupDT_i) with inhibition by perchlorate (p)

$$RupDT _i = \frac{V \max_{DTi \times CT _i} }{Km_{DTi} \times \left(1 + \frac{CT - p}{Km_{DTp}}\right) + CT_i}$$

Allometric Scaling

To account for parameter differences due to varying bodyweights of rats and humans, allometric scaling was applied to maximum velocities (Vmax), PAs, clearance values (Cl), tissue volumes (V) and blood flows (Q). The scaling equations are provided in Merrill, 2000.

RESULTS AND DISCUSSION

Parameterization of Iodide Model

Development of the iodide model was performed by fitting the model to ¹³¹Γ kinetic data from Hays and Solomon (1965). Kms and partition coefficients were available from literature and/or derived by fitting the model to in-house time course data measured in the male rat, as described earlier. Vmaxc values and PAs were derived by fitting the model simulations to radioiodide (¹³¹Γ) uptake in the thyroid and gastric juice, as described in the following paragraphs. The urinary clearance value was derived by fitting model simulations of cumulative iodide in urine to the observed measurements.

In the 1965 study by Hays and Solomon, aspirated gastric juice accounted for an average of 23% of the iv dose within 3 hours of the injection. These data were used to develop the rate of $^{131}\Gamma$ transfer in and out of gastric juice. Simulation of the gastric juice removed during the aspiration session (Figure 2C) required mathematically removing the amount of $^{131}\Gamma$ reabsorbed by the stomach wall. This was accomplished by adjusting the rate of reabsorption of $^{131}\Gamma$ from gastric juice to gastric tissue during the aspiration session. Therefore the term $RAGJ_i$ (the rate of change in $^{131}\Gamma$ in the gastric juice), as it is used in the calculation of RAG_i (the rate of change in $^{131}\Gamma$ in the gastric tissue), described below, is adjusted as follows:

$$RAG_i = PAG_i \times \left(\begin{array}{c} CVGB_i - CG_i \\ -PG_i \end{array} \right) + RAGJ_i - RupGJ_i$$

where:

 $RupGJ_i$ = the rate of active uptake of ¹³¹ Γ at the symporter and secretion into gastric juice and $RAGJ_i$ is calculated under normal conditions (control session) as:

$$RAGJ_i = PAGJ_i \times \begin{pmatrix} CGJ_i / PGJ_i - CG_i \end{pmatrix} + RupGJ_i$$

To remove the secreted $^{131}\Gamma$ in gastric juice from recirculation during the aspiration session, the above equation is modified to eliminate the partitioning of $^{131}\Gamma$ from gastric juice back into systemic circulation as shown below:

$$RAGJ_i = PAGJ_i * (-CG_i) + RupGJ_i$$

The Vmaxc's for the stomach and thyroid were then obtained by fitting values of ¹³¹ Γ uptake into gastric juice from the aspiration session (lower lines in Figures 2B and C). $PAGJc_i$, representing ¹³¹ Γ transfer from the gastric juice into the gastrointestinal plasma was fit to the curve of total ¹³¹ Γ in the aspirated gastric secretions (Figure 2C). The urinary clearance value ($Cluc_i$) was fitted to simulate both cumulative urine content and plasma iodide from the aspiration session data (lower lines in Figures 2A and D).

Once parameters were established using the aspiration session, PAG_i was fitted to predict the corresponding increase in $^{131}\Gamma$ in plasma, thyroid and urine seen in the control session versus the aspiration session. (upper lines in Figures 2A, B and D).

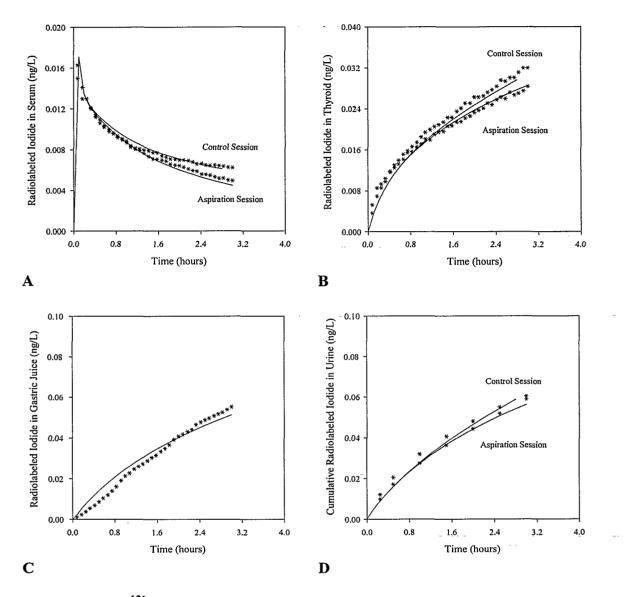


Figure 2. Mean $^{131}\Gamma$ in plasma (A), thyroid (B), gastric juice (C) and urine (D) of nine healthy males after an iv dose of 10 μ Ci $^{131}\Gamma$ (~3.44 ng/kg) (Hays and Solomon, 1965). Model predicted (lines) and actual values (stars) are presented for both the control and aspiration sessions.

Kinetic data on radioiodide or perchlorate uptake in human skin were not available. Effective partitions, described above, were estimated from literature (Table 2). A saturable skin compartment was required in order to fit the human serum ¹³¹Γ concentrations in Figure 2. In addition, the trend of the simulated human skin concentrations was compared to the trend seen in the male rat time course data (see corresponding male rate model in Merrill (2000)). In the male rat, skin ¹²⁵Γ concentrations plateau approximately 2.3 times higher than serum concentrations within 4 hour after injection. ¹²⁵Γ appears to diffuse slowly out of skin. Therefore, after

Vmaxc_S and PASkc values were adjusted to achieve serum fits, the simulated relationship between skin and serum iodide concentrations were compared with those of the male rat (Figure 3 B and D).

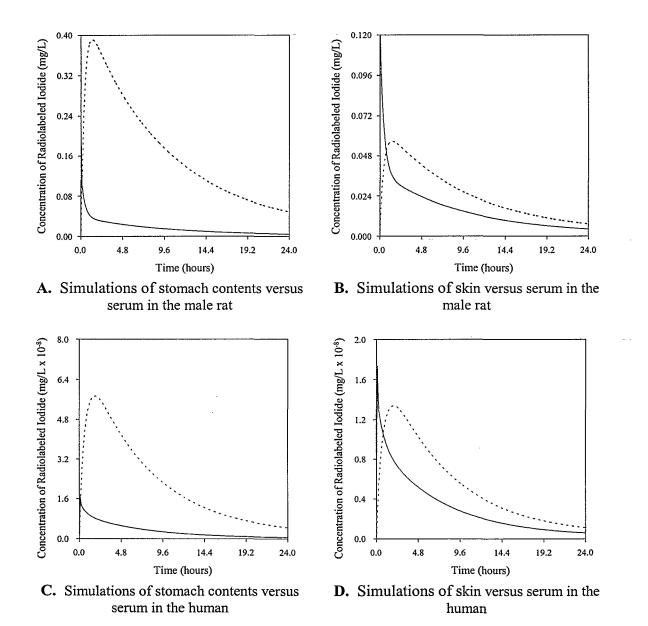
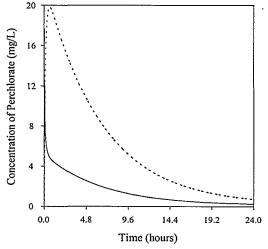


Figure 3. Comparison of model simulations of ¹³¹ I concentrations in stomach contents (dotted line) and skin (dashed line) in relation to plasma ¹³¹ I concentration (solid line) in the male rat (A and B) and the human (C and D). Corresponding doses between the male rat and human are not shown. The simulations in the male rat (A and B) represent actual fits from an *iv* dose of 0.033 mg/kg ¹²⁵ I taken from Merrill (2000). In the human simulations (C), plasma and stomach contents are actual fits from an *iv* dose of 3.44 ng/kg ¹³¹ I (see Figures 2 A and C), while the skin simulation (D) is hypothetical.

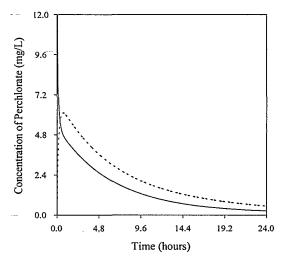
Parameterization of the Perchlorate Model

As with the iodide model, partitions and Km values were derived from the literature and in-house studies, leaving Vmax values and PAs to be derived from model fitting. Kinetic perchlorate data in humans were limited to serum and urine concentrations. Therefore, predicted perchlorate levels in human stomach contents, skin and thyroid are hypothetical. The fitting of the iodide model described above resulted in parameters similar to those of the male rat (Merrill, 2000) with the exception of the Vmaxc values of the thyroid follicular epithelium and colloid. Based on this comparison, it was assumed that perchlorate parameters in the human would also share similar values to those fitted for the rat.

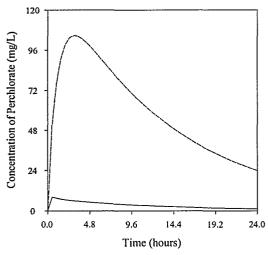
PAs and Vmax values of the stomach and skin were adjusted to simultaneously simulate serum perchlorate concentrations and cumulative perchlorate excreted in urine, established by adjusting the urinary clearance value (ClU_p). Resulting trends of the predicted human stomach and skin levels over time were then compared to the same trends in the male rat and found to be similar (Figure 4 A-D).

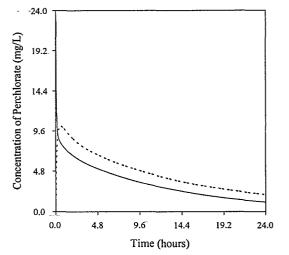


A. Simulated stomach contents (dotted lines) and serum (solid line) in male rat



B. Simulated skin (dotted lines) and serum (solid line) in male rat





C. Simulated stomach contents (dashed lines) and serum (solid line) in human

D. Simulated skin (dashed lines) and serum (solid line) in human

Figure 4. Comparison of model simulations of ClO₄ concentration in stomach contents and skin (dotted line) in relation to plasma ClO₄ concentration (solid line) in the male rat (A and B) and human (C and D) after an *iv* dose of 3 mg/kg ClO₄. Rat simulations (A and B) are based on actual fits of data taken from Merrill (2001). Human simulations (C and D) are hypothetical fits.

Plasma Protein Binding

Early model simulations at 0.1 mg/kg-day underestimated serum perchlorate concentrations while still fitting urinary excretion of ClO₄⁻ (Figure 5A and 6). The low serum predictions suggested either less uptake into other tissues or plasma protein binding. Because serum concentrations in the 0.5 mg/kg-day dose group were fitted using the same parameters, plasma protein binding appeared to be the most likely hypothesis. Increasing uptake in other tissues such as skin and gut resulted in the underestimation of serum levels at 0.5 mg/kg-day. In addition, literature studies suggest perchlorate binding in human serum (Scatchard and Black,1949; Hays and Green, 1973). Plasma binding was modeled to successfully achieve fits of model simulations to serum ClO₄⁻ data at the 0.5, 0.1 and 0.02 dose groups with a common set of parameters (Figure 5B). The model indicates that humans have a lower binding capacity for ClO₄⁻ than rats. For example the Vmaxc for ClO₄⁻ is 9.3x10³ ng/hour-kg in the male rat versus 5.0x10² ng/hour-kg in the human.

The literature suggests that the hypothesized binding site for perchlorate is albumin. A discussion of these studies is provided in Merrill (2001). The difference between adding plasma binding and not is subtle at 0.5 mg/kg-day; however binding does improve uptake and clearance fits in the 0.1 and 0.02 mg/kg-day groups.

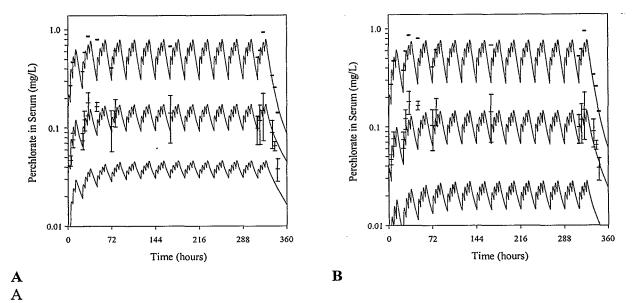


Figure 5. Model simulations (lines) with plasma binding (A) and without (B) and means and standard deviations in the observed values (cross bars) from 4 male subjects dosed 0.02, 0.1 and 0.5 mg/kg-day for 14 days (Greer *et al.*, 2000).

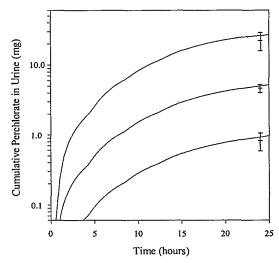
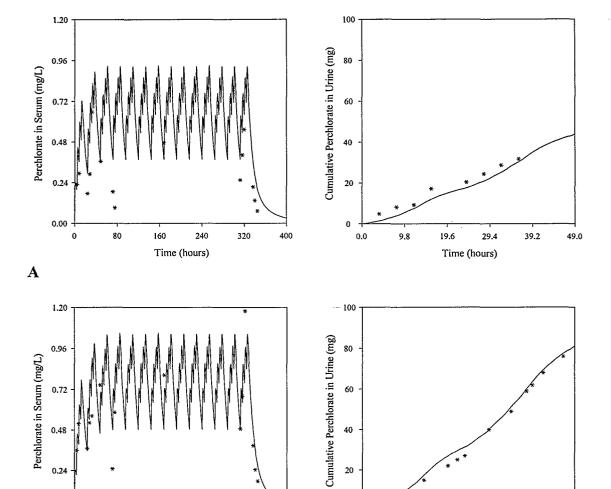


Figure 6. Model simulations (lines) and mean and standard deviations of the observed cumulative urine (cross bars) in male subjects dosed 0.02, 0.1 and 0.5 mg/kg-day (Greer *et al.*, 2000)

Serum and cumulative urine perchlorate levels were simulated for each individual in the 0.5, 0.1 and 0.02 mg/kg-day dose groups. An average value for urinary clearance of perchlorate, ClUc_p, of 0.126 L/hour-kg (± 0.050) was calculated from the individually fitted values. The following plots show the fit obtained for several subjects obtained by using the average value for ClUc_p (Figures 7 through 9).



0.0

400

9.8

19.6

29.4

Time (hours)

39.2

49.0

В

0.00

0

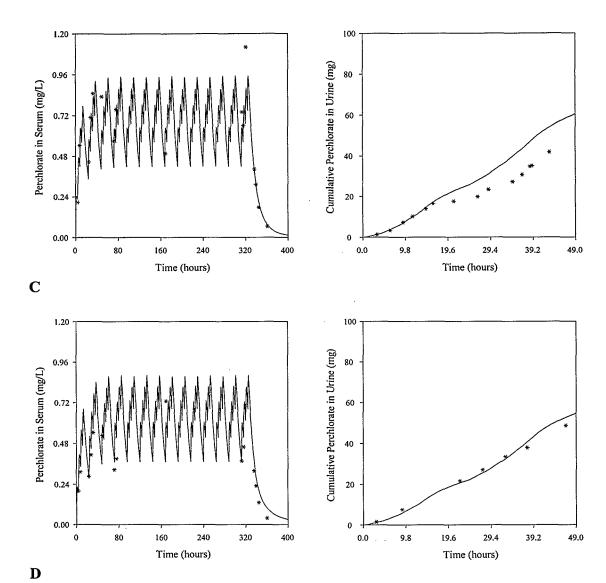
80

240

320

160

Time (hours)



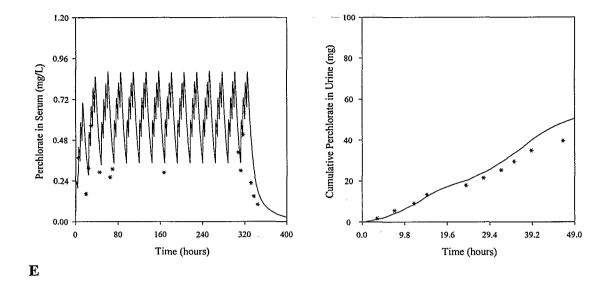
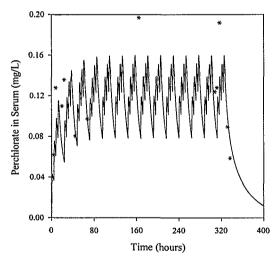
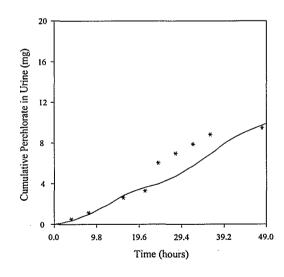
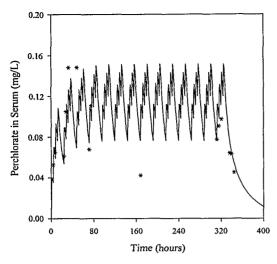


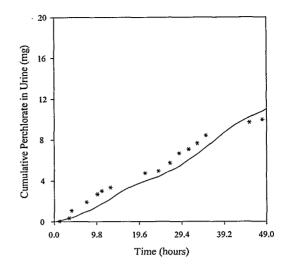
Figure 7. Model predicted (lines) and actual values (asterisks) of individuals' serum perchlorate concentrations and their corresponding 48 hour cumulative urine perchlorate amounts from 4 healthy subjects who consumed 0.5 mg/kg-day perchlorate in drinking water, 4 times per day for 14 days. Model predictions obtained by using the average value for ClUc_p (Greer et al., 2000).



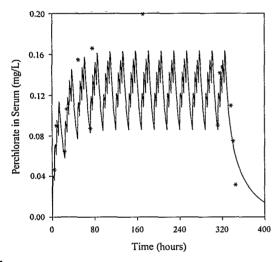


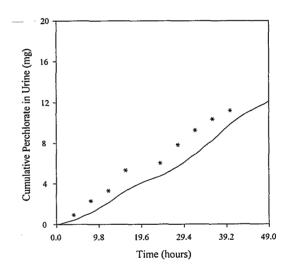
A



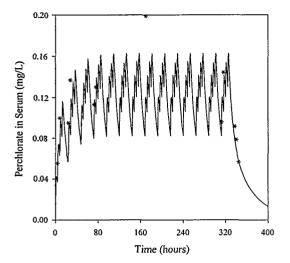


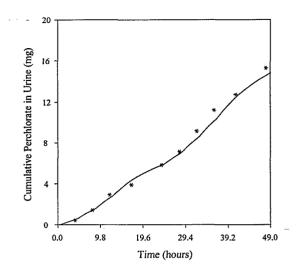
В





 \mathbf{C}

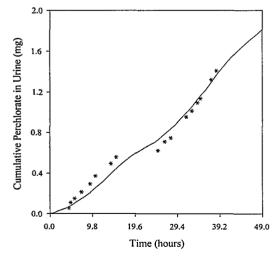


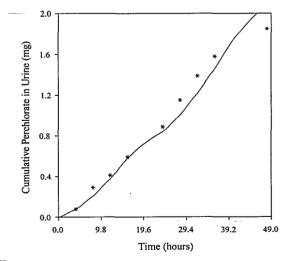


D

Figure 8. Model predicted (lines) and actual values (asterisks) of individuals' serum perchlorate concentrations and their corresponding 48-hour cumulative urine perchlorate from 4 healthy subjects who consumed 0.1 mg/kg-day perchlorate in drinking water 4 times per day for 14 days. Model predictions obtained by using an average ClUc_p (Greer et al., 2000).

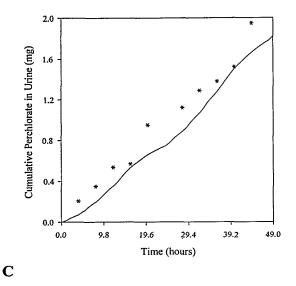
Serum perchlorate levels were not available for the 0.02 mg/kg-day dose group; however, cumulative urine values were fitted (Figure 9) using the average ClUc_p (0.126 L/hour-kg) calculated from the individual fits in the 0.1 and 0.5 mg/kg-day dose groups.





A

 \mathbf{B}



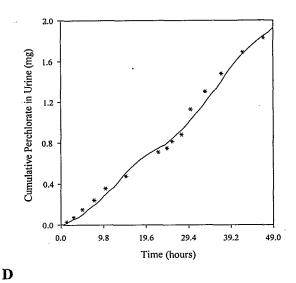


Figure 9. Model simulations (lines) and experimental data (asterisks) of cumulative perchlorate in urine over 48 hours from 4 healthy subjects who consumed 0.02 mg/kg-day perchlorate in drinking water. Model predictions obtained by using the average ClUc_p (Greer et al., 2000).

Perchlorate Induced Inhibition of Thyroid Iodide Uptake

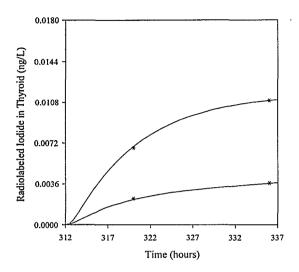
Due to its small size, variations in the thyroid parameters have little effect on serum concentrations of both iodide and perchlorate. As described previously, iodide parameters, including those of the thyroid, were estimated from fits of the data from Hays and Solomon (1965)(see Table 2 and Figure 2B). Using the same iodide parameters, baseline thyroid radioiodide uptakes were fitted by adjusting the Vmax for the follicular epithelium (Vmaxc_Ti) (Figures 10 through 14). An average Vmaxc_Ti (1.5x 10⁵ ng/hour-kg) was obtained from fitting baseline radioiodide uptake measurements provided by Greer *et al.* (2000) across doses (see Table 3). The large variability in Vmaxc_Ti, ranging from 5.0x10⁴ to 5.0x10⁵ ng/hour-kg, may be attributed to variability in endogenous iodide levels, as dietary iodide was not controlled. All participants in each dose group were healthy and screened for thyroid disorders.

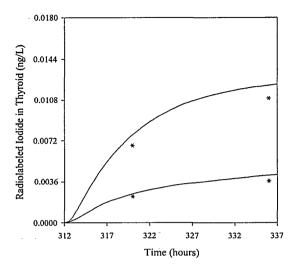
Thyroid parameters for perchlorate were assumed to be similar to those of the male rat, with exception of the Vmax value in the follicular epithelium and colloid (Vmaxc_Tp and Vmaxc_DTp, respectively), which were increased approximately by the same proportional increase in Vmaxc_Ti and Vmaxc_Dti from the male rat to the human (established by fitting the thyroid radioiodide data by Hays and Solomon (1965)). Table 3 summarizes the individually fit urinary clearance constants for perchlorate and Vmaxc_Ti uptake resulting from each subject in the Greer *et al.* (2000) study.

Table 3. Individually fitted ClUc_p(s) and Vmaxc_Ti(s) (Greer et al., 2000)

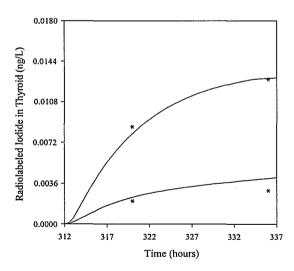
			ClUc_p	Vmaxc_Ti
Dose Group	Subject	Sex	(L/hr-kg)	(ng/hr/kg)
0.5 mg/kg/d	AN	f	0.10	1.34E+05
_	DR	m	0.09	1.20E+05
	JS1	m	0.10	2.00E+05
	CW	f	0.09	1.80E+05
	TO	m	0.09	2.50E+05
	MA	m	0.09	8.00E+04
	AB1	f	0.10	5.00E+05
	RC	f	0.10	1.00E+05
Group Mean			0.10	1.96E+05
Group Std Dev			0.01	1.35E+05
0.1 mg/kg/d	RT	f	0.10	1.10E+05
	NR	m	0.20	2.20E+05
	KN	m	0.20	6.80E+04
	JF	\mathbf{f}	0.12	1.50E+05
	RB1	m	0.24	1.20E+05
	AH	\mathbf{f}	0.10	1.60E+05
	SG	\mathbf{f}	0.13	5.00E+04
	AB2	m	0.17	1.20E+05
Group Mean			0.16	1.25E+05
Group Std Dev			0.05	5.35E+04
0.02 mg/kg/d	SV	f	NA	1.50E+05
	CB	f	0.10_{-}	1.40E+05
	QY	m	0.15	8.00E+04
	DH	m	*	1.50E+05
	JS2	m	NA	1.40E+05
	SK	f	0.20	1.50E+05
	DC	f	0.11	9.00E+04
	GB	m	0.06	8.00E+04
Group Mean			0.12	1.23E+05
Group Std Dev			0.05	3.28E+04
0.007 mg/kg/d	RB	f	NA	1.35E+05
	PE	m	NA	1.35E+05
	MJ	\mathbf{f}	NA	7.80E+04
	SE	\mathbf{f}	NA	8.00E+04
	EA	\mathbf{f}	NA	1.40E+05
	LB	\mathbf{f}	NA	2.80E+05
	LR	\mathbf{f}	NA	9.00E+04_
Group Mean			NA	1.34E+05
Group Std Dev			NA	6.99E+04
Total Mean	7-8-3		0.13	1.45E+05
Total Std Dev			0.05	8.17E+04

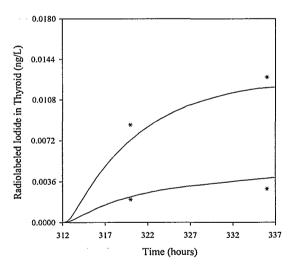
Note: The 0.007 mg/kg-day dose was run to obtain inhibition data only. Serum and urine samples were not collected.



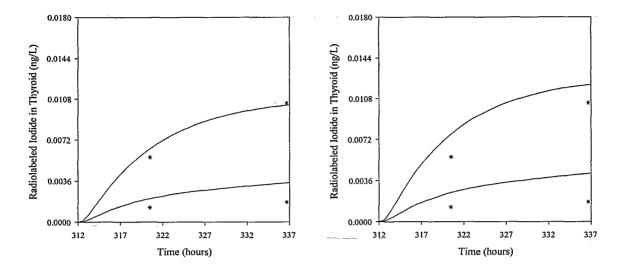


A. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.5 mg/kg-day (bottom asterisks) in a healthy female. Simulation on left obtained by using individually fitted Vmaxc_Ti of 1.3x10⁵. Right simulation obtained using average Vmaxc_Ti.

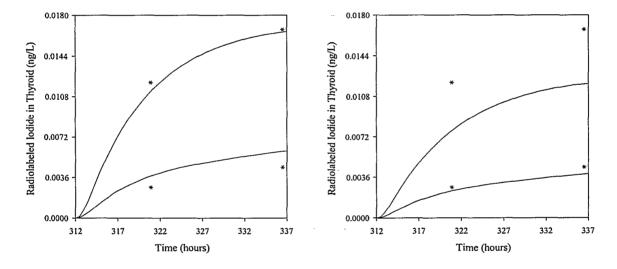




B. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.5 mg/kg-day (bottom asterisks) in a healthy female. Simulation on left obtained by using individually fitted Vmaxc_Ti of 1.8x10⁵. Right simulation obtained using average Vmaxc_Ti.

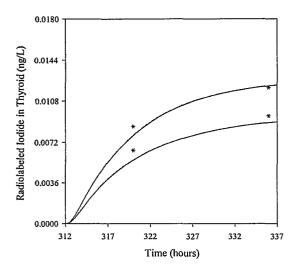


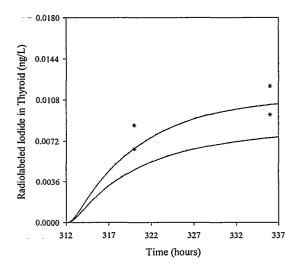
C. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.5 mg/kg-day (bottom asterisks) in a healthy male. Simulation on left obtained by using individually fitted Vmaxc_Ti of 1.24x10⁵. Right simulation obtained using average Vmaxc_Ti.



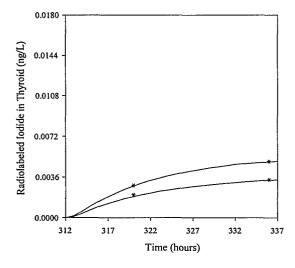
D. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.5 mg/kg-day (bottom asterisks) in a healthy male. Simulation on left obtained by using individually fitted Vmaxc_Ti of 2.5x10⁵. Right simulation obtained using average Vmaxc_Ti.

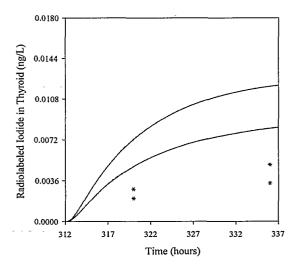
Figure 10. Model predicted (lines) and actual 8 and 24 hour RAIU measurements (asterisks) from 4 healthy subjects before perchlorate exposure (upper lines and asterisks) and on day 14 of perchlorate exposure at 0.5 mg/kg-day (lower lines and asterisks). Simulations in the figures on the right represent individual fits of thyroid iodide uptake by adjusting Vmaxc_Ti. Simulations in the left figures were established by using the average Vmaxc Ti (Greer et al., 2000).



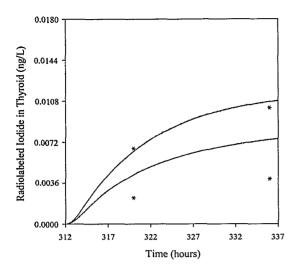


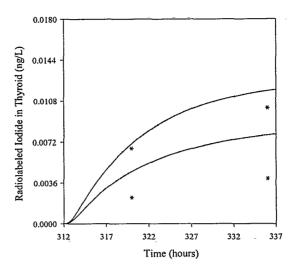
A. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.1 mg/kg-day (bottom asterisks) in a healthy female. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 1.65x10⁵. Right simulation obtained using average Vmaxc_Ti.



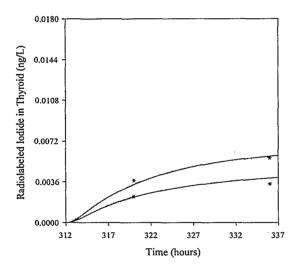


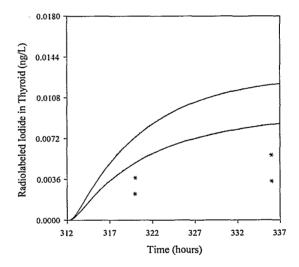
B. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.1 mg/kg-day (bottom asterisks) in a healthy female. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 5.0×10^4 . Right simulation obtained using average Vmaxc_Ti.





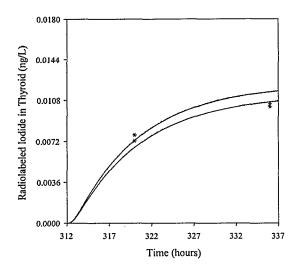
C. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.1 mg/kg-day (bottom asterisks) in a healthy male. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 1.2x10⁵. Right simulation obtained using average Vmaxc_Ti.

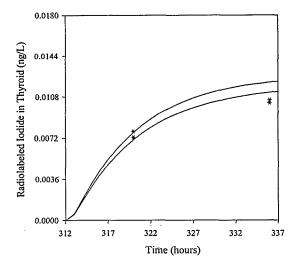




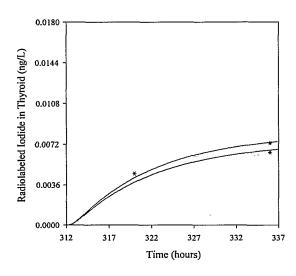
D. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.1 mg/kg-day (bottom asterisks) in a healthy male. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 6.8x10⁴. Right simulation obtained using average Vmaxc_Ti.

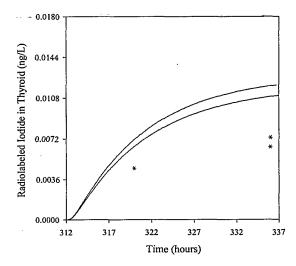
Figure 11. Model predicted (lines) and observed 8 and 24 hour RAIU measurements (asterisks) during baseline (upper lines and asterisks) and on day 14 of perchlorate exposure at 0.1 mg/kg-day (lower lines and asterisks) from 4 healthy subjects. Simulations on the left were obtained by individual fits of thyroid radioiodide uptakes by adjusting Vmaxc_Ti. Simulations on the right were obtained by using an average Vmaxc_Ti of 150,000 ng/hour-kg (Greer et al., 2000).



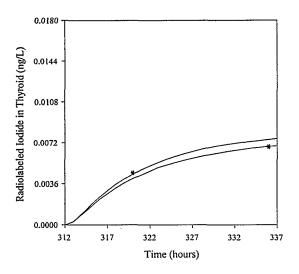


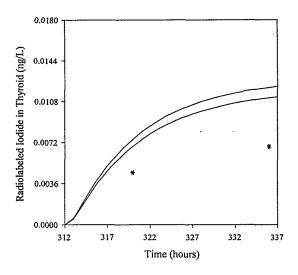
A. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.007 mg/kg-day (bottom asterisks) in a healthy female. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 1.4x10⁵. Right simulation obtained using average Vmaxc_Ti.



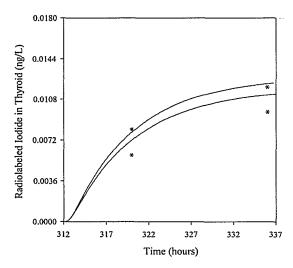


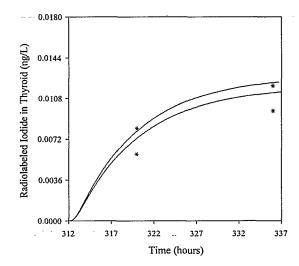
B. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.02 mg/kg-day (bottom asterisks) in a healthy male. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 8.0x10⁴. Right simulation obtained using average Vmaxc_Ti.





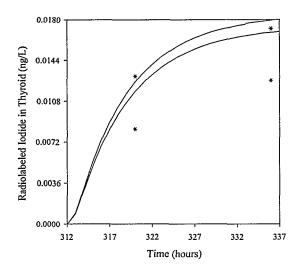
C. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.02 mg/kg-day (bottom asterisks) in a healthy male. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 8.0x10⁴. Right simulation obtained using average Vmaxc_Ti.

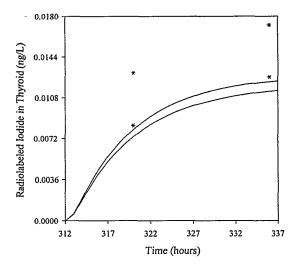




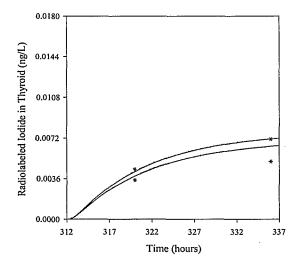
D. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.02 mg/kg-day (bottom asterisks) in a healthy female. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 1.5x10⁵. Right simulation obtained using average Vmaxc_Ti.

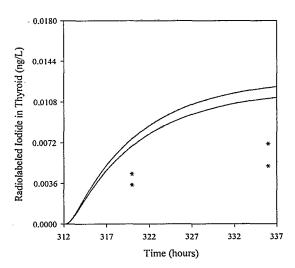
Figure 13. Model predicted (lines) and observed 8 and 24 hour RAIU measurements (asterisks) during baseline (upper lines and asterisks) and on day 14 of perchlorate exposure at 0.02 mg/kg-day (lower lines and asterisks) from 4 healthy subjects. Individually fit thyroid uptakes for each subject are displayed in the plots on the left. Simulated thyroid uptakes using an average Vmaxc_ti of 150000 ng/hour-kg are displayed in the right plots (Greer et al., 2000).



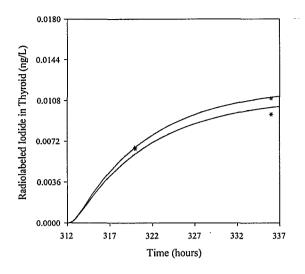


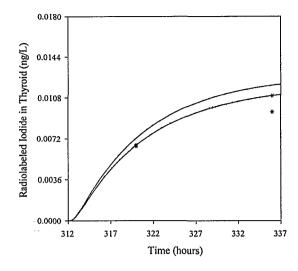
A. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.007 mg/kg-day (bottom asterisks) in a healthy female. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 2.8x10⁵. Right simulation obtained using average Vmaxc_Ti.



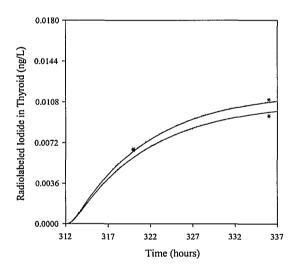


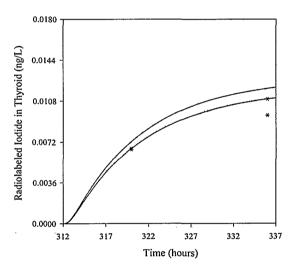
B. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.007 mg/kg-day (bottom asterisks) in a healthy female. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 7.8x10⁴. Right simulation obtained using average Vmaxc_Ti.





C. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.007 mg/kg-day (bottom asterisks) in a healthy male. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 1.35x10⁵. Right simulation obtained using average Vmaxc_Ti.





D. Thyroid radioiodide uptakes (before perchlorate) (top asterisks) and on day 14 of perchlorate exposure at 0.007 mg/kg-day (bottom asterisks) in a healthy female. Simulation on the left obtained by using individually fitted Vmaxc_Ti of 1.35x10⁵. Right simulation obtained using average Vmaxc_Ti.

Figure 14. Model predicted (lines) and observed 8 and 24 hour RAIU measurements (asterisks) during baseline (upper lines and asterisks) and on day 14 of perchlorate exposure at 0.007 mg/kg-day (lower lines and asterisks) from 4 healthy subjects. Individually fit thyroid uptakes for each subject are displayed in the left plots. Simulated thyroid uptakes using an average Vmaxc_ti of 150000 ng/hour-kg are displayed in the right plots (Greer et al., 2000).

The average Vmaxc_Ti fitted from Greer et al. (2000) (1.5 x 10⁵ ng/hour-kg) agrees well with the value obtained from fitting data from Hays and Solomon (1965) (2.5 x 10⁵ ng/hour-kg). The subjects in the Hays and Solomon study had fasted 12 hours prior to the administration of the *iv* radioiodide dose. As a result, intrathyroidal iodide levels would have been lower in the fasted individuals than in the individuals in the study by Greer et al. (2000), in which no dietary restrictions prior to thyroid uptake measurements were implemented. Therefore, the average Vmaxc_Ti from Hays and Solomon (1965) is expected to exceed that obtained from Greer et al. (2000).

Table 4 lists the average serum perchlorate concentrations and percent inhibitions from exposure day 14 from each dose group in the Greer *et al.* (2000) study described above. In addition, the average value for serum perchlorate concentration is provided from the unpublished study by Drs. Brabant and Leitoff, described previously in the Methods section. Simulations of data from this study are provided in the Model Validation section.

Table 4. Average serum perchlorate and percent inhibition across dose groups in the 14day drinking water studies

Dose (mg/kg-day)	BW (kg)	Serum ClO4 ⁴ (mg/L)	Average % Inhibition of 24 hour Iodide Uptake ⁴	Data Source
				Unpublished data from
12.0	83.5 ± 17.4	23.0 ± 13.2	NA^{1}	Brabant and Leitolf, 2000
0.5	76.1 ± 14.3	0.649 ± 0.25	67.4 ± 12.1	Greer et al., 2000
0.1	78.7 ± 13.2	$0.126 \pm .053$	43.4 ± 12.3	Greer et al., 2000
0.02	78.4 ± 18.2	<DL ²	18.2 ± 12.8	Greer et al., 2000
0.007	73.2 ± 15.4	NA^3	6.0 ± 22.0	Greer et al., 2000

NA = not available

MODEL VALIDATION

The model was developed using the iodide kinetic data from Hays and Solomon (1965) and serum and urine perchlorate data with radioiodide uptake and inhibition measurements in the thyroid provided by Greer and associates (2000). The ability of the model to predict human data

¹RAIU measurements were not taken in the unpublished 12 mg/kg/d study by Brabant and Leitolf.

² Samples at or below detection limit

³The 0.007 mg/kg/d dose was run to obtain inhibition data only. Serum and urine samples were not collected.

⁴Averages provided were calculated from exposure day 14 only.

from other experiments, analyzed at different labs, was tested using available data from independent studies.

Using the parameters in Table 2, the model adequately simulated cumulative perchlorate in urine reported in three published studies using therapeutic perchlorate dose levels (Figures 15 through 17). Oral doses administered in these studies were approximately 9.07 mg/kg (Durand, 1938), 9.56 mg/kg (Kamm and Drescher, 1973) and 20 mg/kg (Eichler, 1929). The previously determined urinary clearance value (ClUc_p) of 0.126 L/hour-kg was used with all validation data.

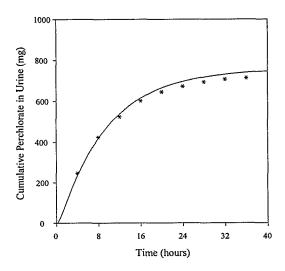


Figure 15. Model predicted (line) and observed (circles) cumulative ClO₄ in urine from a healthy male after an oral dose of 9.56 mg ClO₄ (Kamm and Drescher, 1973). Simulation obtained by using parameters in Tables 1 and 2.

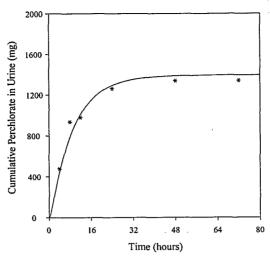


Figure 16. Model predicted (line) and observed (circles) cumulative ClO₄ in urine from a healthy male after an oral dose of approximately 20 mg ClO₄/kg (Eichler, 1929). Simulation was obtained by using the parameters in Tables 1 and 2.

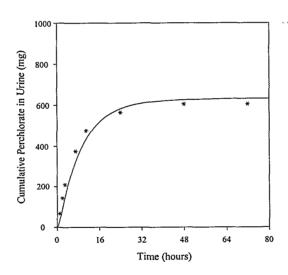
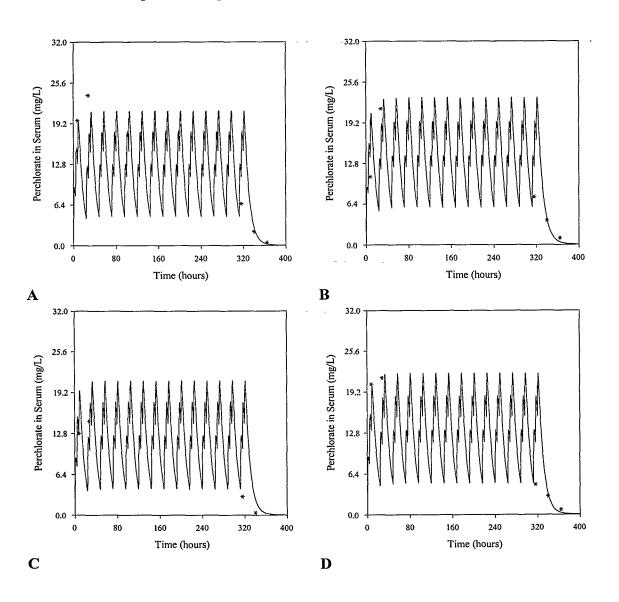


Figure 17. Model predicted (line) and observed (circles) cumulative amount of ClO₄⁻ in urine from a healthy male after an oral dose of approximately 9.07 mg/kg ClO₄⁻ (Durand, 1938). Simulation obtained by using the parameters in Tables 1 and 2.

The ability of the model to predict cumulative perchlorate in urine from three different studies at three different doses with the same set of parameters, established from the studies by Hays and Solomon (1965) and Greer *et al.* (2000), demonstrates the usefulness of the model. In addition is provides validation for the model structure and physiological and chemical parameters used.

The model also predicts serum perchlorate concentrations at 12 mg/kg-day from an unpublished study performed by Dr. Georg Brabant at the Medizinische Hochschule, Hanover, Germany (Figure 18). This study was very similar to Greer *et al.* (2000). Subjects received 12 mg/kg-day perchlorate in drinking water near meal times. Variability in the observed serum measurements is believed to reflect variability in the dosing regime, as the experimental protocol was less fixed than that used in Greer *et al.* (2000). Again the usefulness of the model is demonstrated by its ability to successfully predict serum concentrations from a dose 24 times higher than the high dose used to establish perchlorate parameters (0.5 mg/kg-day).



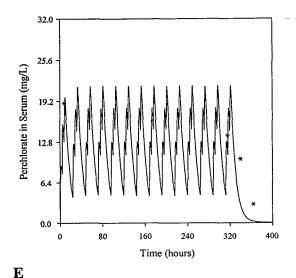


Figure 18. Serum perchlorate concentrations in 5 males during exposure to 12 mg/kg-day ClO₄ in drinking water. Subjects were instructed to ingest the drinking water solution 3 times/day for 14 days. Serum samples were collected at approximately 2 hours after the first dose, after 12 pm on day two, the morning of day 14 and post exposure days 1 and 2 (unpublished data, Brabant and Leitolf).

The model successfully simulates the thyroidal iodide uptake in a subject with hyperthyroidism by increasing the maximum velocity in the follicular epithelium (Vmaxc_Ti) to 5.0E6 ng/L-kg, a factor of ten times higher than in normal subjects (upper line in Figure 19). This increase in Vmaxc_Ti is supported in the literature, as Gluzman and Niepomniszcze (1983) measured elevated Vmax(s) in thyroid specimens from subjects with Grave's disease. That the model is capable of predicting uptake under this condition supports the usefulness of the current model structure.

The model underpredicts the degree of inhibition caused by perchlorate in this subject (bottom line Figure 19). It would appear that the increased inhibition could be attributed to a lower Km value. However, Gluzman and Niepomniszcze (1983) noted that the Km did not differ greatly between thyroid specimens from hyperthyroid subjects and normal subject. This suggests that the increased inhibition by perchlorate seen in Grave's disease may be attributed to a mechanism other than NIS affinity.

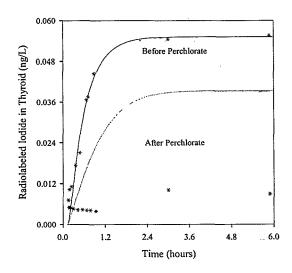


Figure 19. Model predicted (line) and observed (asterisks) amount of ¹³¹Γ uptake in the thyroid of a male with Graves after *iv* dose of 10 μCi ¹³¹Γ before and after 100 mg KClO₄ (Stanbury and Wyngaarden, 1952).

SUMMARY AND CONCLUSIONS

The model adequately simulates serum concentrations and cumulative urine after drinking water exposure to perchlorate spanning four orders of magnitude (0.02 to 12.0 mg/kg-day). Serum perchlorate levels were not available at 0.02 mg/kg-day; however the model did predict the cumulative urine from that dose group (Figure 10). The model slightly underpredicts serum levels at 0.1 mg/kg-day (Figure 9). It is possible that binding with some serum proteins occurs to a greater extent than demonstrated in this model; however, increasing binding does not allow fitting of the urine data at that dose level, so it's possible that less uptake in one or more tissues is actually occurring.

Aspects of the model, which were supported in the literature or laboratory studies but which could not be directly observed in humans, were incorporated if necessary to improve fits. For example, the slow diffusion of iodide and active uptake of perchlorate into human skin was incorporated in spite of a lack of available human data to support them. However, the model required the inclusion of active uptake in human skin. Without the skin compartment, the model overestimated circulating plasma inorganic iodide and perchlorate. As discussed earlier, cutaneous uptake of iodide and perchlorate in mice and rats has been reported (Brown-Grant and Pethes, 1959; Zeghal *et al.*, 1995). The lack of reported iodide in human skin from clinical radioiodide scans may be due to difficulty in differentiating it from background radioactivity. However, due to its large size, skin appears to be an important pool for slow turnover of iodide.

Early iodide kinetics were established by fitting data from Hays and Solomon (1965), as described previously. Interestingly, the simulated amount of ¹³¹ I in gastric juice (simulated control session) indicates rapid uptake of iodide (and likely perchlorate as well) into this

compartment and quick reabsorption in the tissue (Figure 2C). GI clearance of iodide is rapid and plays an important role in radioiodide conservation.

The appearance of time-course radioiodine in stomach contents of any species is complicated by the fact that it reflects more than sequestration of radioiodide by NIS. Its appearance also reflects radioiodide contributed through the gradual accumulation of iodide in saliva that is swallowed involuntarily throughout the study. Several studies that examined sequestration of these anions in digestive juices have all shown high variability in the concentrations measured over time (Honour *et al.*, 1952; Hays and Solomon, 1965; Merrill, 2001). There is a tendency for the gastric juice to plasma ratio to be low when the rate of secretion of juice is high (Honour *et al.*, 1952). Fluctuations in the secretion rate are probably the most important factor in determining the pattern of the concentration ratios in individuals. Therefore, variability in stomach or GI tract parameters between models is expected. However, the early rise in the gastric juice:plasma ratio mentioned earlier is a constant feature across these data sets, whether or not an attempt was made to eliminate contamination of gastric juices by dietary contents or saliva. The model successfully predicts this same trend.

Variability across data sets was noted in the mean Vmaxc_Ti values, ranging from 5.0E4 to 5.0E5 ng/hour-kg. These values were estimated from best visual fits of baseline 8 and 24 hour thyroid RAIU data. Inhibition data from higher dose groups would be useful to test the robustness of the model to predict inhibition of uptake of iodide in the thyroid.

Average urinary clearance values were found to be 0.1 L/hour-kg for iodide and 0.126 L/hour-kg for perchlorate. Excretion constants were highest among the 0.1 mg/kg-day group. With the urinary excretion rates fitted to cumulative urine data, the model tends to slightly underestimate serum perchlorate levels at repeated low doses. Elevated serum concentrations may indicate plasma binding of perchlorate. Yamada and Jones (1967) studied effects of different anions on plasma binding to thyroxine and noted that some of the thyroxine had been displaced after perchlorate was introduced. Thus, it is possible that perchlorate competes with thyroxine for the same binding sites of plasma proteins (Merrill, 2001; Clewell, 2001a).

Dietary iodine and endogenous inorganic iodide levels are clearly important in modeling iodide and perchlorate kinetics, because excessive iodide levels cause the ion to inhibit its own uptake. Plasma inorganic iodide (PII) is rarely reported in the literature, due to analytical difficulties, and it was not available in any of the studies presented in this paper. While measurements of tracer radioiodide can be fitted to predict transfer rates, its use is limited when attempting to predict the saturation of nonlinear compartments, such as the thyroid, which are dependent upon the existing amount of iodide already present. Subsequent modeling efforts on predicting subsequent effects of iodide inhibition on thyroid hormone synthesis and regulation in humans will require the capability of the model to predict PII. Ultimately, regional dietary iodine should be considered in creating recommendations on drinking water levels for perchlorate.

Statistical analyses were performed on the hormone data collected from Greer *et al.* (2000). Details on the statistical analyses are provided in Attachment 2. In summary, there was little effect of perchlorate on levels of T₄, free T₄ or T₃. TSH decreased significantly from baseline by

exposure day 3. On post exposure day 1, the TSH levels of the subjects in the 0.5 mg/kg-day group had decreased by an average of 35% from baseline (ranging from 17% to 52%). Therefore, it appears that TSH was dropping while inhibition remained the same. It is possible that there is an increase in thyroid sensitivity to TSH as an early response to inhibition (Brabant et al., 1992). This increased sensitivity (possibly an increase affinity of the TSH receptor) could possibly decrease circulating TSH levels, while T₄ has not decreased sufficiently yet to stimulate the hypothalamus to increased TRH secretions. After perchlorate was discontinued, between post exposure days 1 and 15, the mean TSH level increased significantly over baseline (23% greater than baseline), with TSH of one subject remaining below baseline. The drop in TSH during perchlorate exposure and the rise above baseline measurements after perchlorate are the opposite of the TSH regulation expected and are unexplained at this point.

In addition, the data by Greer *et al.* (2000) showed an increase in radioiodide uptake in excess of baseline measurements 14 days after perchlorate exposure. An increase in radioiodide uptake is expected due to the rise in TSH mentioned above. This rebound effect has been noted in other human inhibition studies (using both iodide and perchlorate as inhibitors). Saxena *et al.* (1962) evaluated the prophylactic doses of iodide required to suppress thyroid uptake of $^{131}\Gamma$ in euthyroid mentally defective children. They found a minimal effective oral dose of 1500 to 2000 µg iodide per square meter of body surface per day was required to completely suppress $^{131}\Gamma$ uptake. Within a week after iodide administration was stopped, a rebound of uptake was noted. In some instances these uptakes were even higher in subsequent weeks.

The PBPK models developed for perchlorate induced inhibition may ultimately be used to evaluate the dose-response of adverse effects from low level perchlorate exposure. Modeled effects on hormone regulation are yet to be developed. Perturbations in hormones levels after perchlorate exposure demonstrate complex differences in the hormone regulatory mechanisms between rats and humans, which are difficult to describe (Merrill, 2001; Clewell, 2001a and 2001b). However, the current model structures may also provide a basis for evaluating thyroid effects from other environmental contaminants. For example, excessive exposure to other similarly behaving anions, such as nitrate (a frequently found contaminant in drinking water supplies), may contribute to antithyroid effects (Wolff and Maurey, 1963).

ACKNOWLEDGEMENTS

The authors express special thanks to Drs. Monte Greer, Gay Goodman, Georg Brabant and Holger Leitolf for providing serum and urine samples from their experiments and sharing their results. Also acknowledged are Lt Col Dan Rogers, Dr. Richard Stotts and Dr. Dave Mattie, U.S. Air Force, for assistance in obtaining funding of this research and Mel Andersen (Colorado State University, Fort Collins, CO), Harvey Clewell (ICF Consulting, Ruston, LA) and Annie Jarabek (NCEA, USEPA, RTP, NC) for expert advice. The authors would also like to acknowledge Charles Goodyear for performing statistical analyses of the data. Mr. Goodyear is a statistical consultant for AFRL, Human Effectiveness Directorate, Crew Systems Interface Division (AFRL/HEC), Wright-Patterson AFB, OH. Lastly, in-house work would not have been possible without analytical support from Lt Eric Eldridge (AFRL/HEST, WPAFB, OH), Latha Narayanan

(GEOCENTERS, Inc, WPAFB, OH), Gerry Buttler (ManTech Environmental Technology, Dayton, OH) and SSgt Paula Todd (AFRL/HEST, WPAFB, OH).

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ATTACHMENT 2

Serum Hormones (TSH, T₃, T₄, fT₄) Statistical Analysis

Charles D. Goodyear

Twenty-four subjects (12 male and 12 female) participated in the 2000 study by Greer and colleagues. Each subject had their hormone levels (TSH, T₄, free T₄ and T₃) determined on a screening day and a baseline day. At this time, the subjects were randomly assigned to one of 3 dose groups (0.02, 0.1 and 0.5 mg/kg-day) such that there were 4 male and 4 female subjects in each group. An additional dose group (0.007 mg/kg-day) was included for inhibition measurement but no serum samples were collected besides preliminary screening samples. Over the next 2 weeks, each subject was dosed with perchlorate, using drinking water, at different times on days 1, 2, 3, 4, 8 and 14. Before drinking perchlorate, blood was drawn to determine their hormone levels. Lastly, blood was drawn 1 and 15 days post exposure. For all subjects, exposure day 1 was 5 days after baseline.

Table 1. Times of blood draws

Exposure	Time	#
Day	Period	Subjects
Baseline	1	24
1	2	23
	3	24
	1	23
2	2	23
	3	24
3	1	24
4	1	23
	2	24
	1	15
8	2	7
	3	2
	1	24
14	2	24
	3	24
Post 1	1	24
Post 15	1	24

Time periods were: 1 = 0800 - 0955 hours, 2 = 1010 - 1405 hours and 3 = 1450 - 1946 hours. The only exception was one Post 1 blood draw at 1024 (listed as time period = 1). The time 1946 hours is most likely the recording time instead of the draw time.

On exposure day 8, each subject had blood drawn only once, at different times throughout the day. The purpose of the analyses was to determine: 1) Do the dose groups differ, at each exposure day separately, in their percent change from baseline? 2) Do the males and females differ, at each exposure day separately, in their percent change from baseline? 3) What combinations of dose group and exposure day show a significant change from baseline. Baseline means and standard deviations of subjects are shown in Table 2.

Table 2. Baseline mean and standard deviation of subjects (N=24)

Hormone	Mean	SD
TSH (μU/mL)	2.16	1.01
T ₄ (μg/dL)	7.34	1.75
free T ₄ (ng/dL)	1.16	0.17
T ₃ (ng/dL)	105.25	15.70

Since blood was drawn at different times during the exposure days, time of day needs to be addressed as a factor, especially since previous research has shown time of day to be an influence (Brabant *et al.*, 1992a and 1992b). A preliminary screening occurred 1 to 34 days before baseline with most subjects screened less than 10 days before baseline. Figure 1 contains the screening day hormone level percent change from the baseline hormone level. All baseline blood draws were between 0800 and 0908. The X-axis values represent the time of day for the screening blood draw. For T₄, free T₄ and T₃, time of day does not appear to be related to hormone level for comparing the screening day and the baseline day. For TSH, most subjects who had blood drawn after 1000 on the screening day showed lower hormone levels compared with the baseline day.

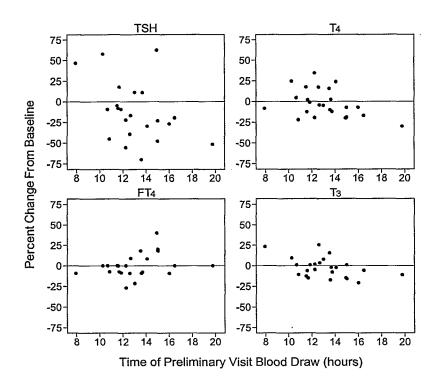


Figure 1. Percent change from baseline to the preliminary visit for the 24 subjects. The time of 1946 is most likely recording time instead of draw time.

Analyses of variance were performed on exposure days 1, 2, 4 and 14, for each dose group separately, to help indicate whether time of day might be related to a change in hormone level. The dependent variable was percent change from baseline. These tests indicated that time of day may influence hormone levels, particularly for TSH. As with the comparisons of screening and baseline days, the TSH level appears to decrease from the morning to the afternoon. This circadian trend in TSH has been documented (Brabant *et al.*, 1992a and 1992b). All further analyses, tables and figures used data from the morning blood draws only (i.e., time period = 1) as the baseline blood draws were in the morning. A thorough analysis of the effects of time of day will not be performed here.

To compare the dose groups and sexes, at each exposure day separately, a two-factor analysis of variance was performed. The dependent variable was the percent change from baseline. Post-hoc paired comparisons of the dose groups used the Bonferroni procedure with a family-wise error level of 0.05. Tables 3 through 6 show results of these analyses. The error degrees of freedom indicate missing data (i.e., DFE = 18 indicates no missing data). For exposure day 8, there was only 15 subjects, 3 female and 2 male in the 0.02 and 0.1 mg/kg-day dose groups, 2 female and 3 male in the 0.5 mg/kg-day dose group.

Two-tailed t-tests, with error pooled across sex but not pooled across dose, were used to determine which combinations of dose group and exposure day showed a significant difference from baseline ($p \le 0.05$). Two-tailed t-tests, without pooled error, were used to determine which

combinations of dose group, exposure day and sex showed a significant difference from baseline $(p \le 0.05)$.

Table 3. Results from two-factor analysis of variance for TSH percent change

Exposure Day	Source	SS	DF	SSE	DFE	F	P
	Dose	1.76E+03	2	6.69E+03	17	2.23	0.1377
2	Sex	1.66E+01	1	6.69E+03	17	0.04	0.8395
	Dose*Sex	1.60E+03	2	6.69E+03	17	2.04	0.1607
	Dose	2.90E+03	2	7.52E+03	18	3.47	0.0532
3	Sex	6.46E-01	1	7.52E+03	18	0.00	0.9691
	Dose*Sex	1.62E+03	2	7.52E+03	18	1.94	0.1726
	Dose	1.40E+03	2	1.47E+04	17	0.81	0.4622
4	Sex	1.99E+03	1	1.47E+04	17	2.29	0.1482
	Dose*Sex	2.70E+03	2	1.47E+04	17	1.56	0.2394
	Dose	3.72E+03	2	3.58E+03	9	4.67	0.0407
8	Sex	3.29E+02	1	3.58E+03	9	0.83	0.3872
	Dose*Sex	5.01E+02	2	3.58E+03	9	0.63	0.5551
	Dose	3.19E+03	2	1.36E+04	15	1.76	0.2061
14	Sex	7.30E+01	1	1.36E+04	15	0.08	0.7804
	Dose*Sex	1.65E+03	2	1.36E+04	15	0.91	0.4239
	Dose	4.69E+03	2	5.85E+03	18	7.22	0.0050
Post 1	Sex	2.66E+03	1	5.85E+03	18	8.18	0.0104
	Dose*Sex	2.73E+03	2	5.85E+03	18	4.19	0.0320
	Dose	6.59E+02	2	1.32E+04	18	0.45	0.6457
Post 15	Sex	8.02E+02	1	1.32E+04	18	1.09	0.3100
	Dose*Sex	6.06E+03	2	1.32E+04	18	4.12	0.0335

For TSH percent change at exposure day 8, there was a significant main effect of dose group (p = 0.0407). Paired comparisons showed a significant difference between the 0.02 (mean = -1%) and 0.5 (mean = -39%) mg/kg-day dose groups.

For TSH percent change at post exposure day 1, there was a significant interaction between dose group and sex (p = 0.0320). Simple main effect tests for each sex separately showed no significant effect of dose group for the males (p = 0.7188) and a significant effect of dose group for the females (p = 0.0057) with the 0.5 dose group (mean = -45%) significantly different from both the 0.02 dose group (mean = -2%) and the 0.1 dose group (mean = 13%). Also at post exposure day 1, there was a significant difference between males and females for the 0.02 dose group (p = 0.0132) (mean: males = -34%, females = -2%) and the 0.1 dose group (p = 0.0363) (mean: males = -27%, females = 13%) but not for the 0.5 dose group (p = 0.5405).

For TSH percent change at post exposure day 15, there was a significant interaction between dose group and sex (p = 0.0335). The only significant simple main effect was between males and females for the 0.1 dose group (p = 0.0064) (mean: males = -16%, females = 40%).

Table 4. Results from two-factor analysis of variance for T₄ percent change

Exposure Day	Source	SS	DF	SSE	DFE	F	P
	Dose	2.43E+02	2	1.06E+03	17	1.96	0.1721
2	Sex	1.81E+02	1	1.06E+03	17	2.92	0.1056
	Dose*Sex	6.67E+01	2	1.06E+03	17	0.54	0.5940
	Dose	2.53E+02	2	5.22E+03	18	0.44	0.6528
3	Sex	3.22E+00	1	5.22E+03	18	0.01	0.9173
	Dose*Sex	1.08E+02	2	5.22E+03	18	0.19	0.8314
	Dose	3.15E+02	2	1.78E+03	17	1.51	0.2491
4 .	Sex	1.23E+01	1	1.78E+03	17	0.12	0.7361
	Dose*Sex	2.12E+02	2	1.78E+03	17	1.01	0.3842
	Dose	7.43E+02	2	2.30E+03	9	1.45	0.2838
8	Sex	4.88E+02	1	2.30E+03	9	1.91	0.2006
	Dose*Sex	3.50E+02	2	2.30E+03	9	0.68	0.5289
	Dose	1.18E+03	2	4.83E+03	17	2.08	0.1554
14	Sex	8.70E+01	1	4.83E+03	17	0.31	0.5871
	Dose*Sex	8.22E+02	2	4.83E+03	17	1.45	0.2628
	Dose	9.73E+02	2	4.65E+03	17	1.78	0.1990
Post 1	Sex	2.47E+02	1	4.65E+03	17	0.90	0.3555
	Dose*Sex	3.29E+02	2	4.65E+03	17	0.60	0.5590
	Dose	6.08E+02	2	4.86E+03	17	1.06	0.3670
Post 15	Sex	1.76E+02	1	4.86E+03	17	0.61	0.4438
	Dose*Sex	8.12E+02	2	4.86E+03	17	1.42	0.2689

Table 5. Results from two-factor analysis of variance for free T_4 percent change

Exposure Day	Source	SS	DF	SSE	DFE	F	P
	Dose	9.73E+01	2	7.90E+03	17	0.10	0.9012
2	Sex	2.60E+01	1	7.90E+03	17	0.06	0.8157
	Dose*Sex	1.55E+03	2	7.90E+03	17	1.66	0.2188
	Dose	6.55E+01	2	5.97E+03	18	0.10	0.9065
3	Sex	1.84E+01	1	5.97E+03	18	0.06	0.8167
	Dose*Sex	5.82E+02	2	5.97E+03	18	0.88	0.4330
	Dose	2.05E+02	2	5.80E+03	17	0.30	0.7447
4	Sex	1.80E+01	1	5.80E+03	17	0.05	0.8212
	Dose*Sex	7.37E+02	2	5.80E+03	17	1.08	0.3617
	Dose	1.22E+02	2	3.45E+03	9	0.16	0.8554
8	Sex	7.72E+01	1	3.45E+03	9	0.20	0.6645
	Dose*Sex	1.76E+02	2	3.45E+03	9	0.23	0.7997
	Dose	4.91E+01	2	6.09E+03	14	0.06	0.9453
14	Sex	3.65E+02	1	6.09E+03	14	0.84	0.3750
	Dose*Sex	2.85E+02	2	6.09E+03	14	0.33	0.7257
	Dose	8.53E+00	2	6.63E+03	18	0.01	0.9885
Post 1	Sex	2.08E+02	_ 1	6.63E+03	18	0.56	0.4626
	Dose*Sex	1.13E+03	2	6.63E+03	18	1.53	0.2439
	Dose	4.16E+02	2	8.40E+03	18	0.45	0.6472
Post 15	Sex	2.83E+02	1	8.40E+03	18	0.61	0.4459
	Dose*Sex	1.54E+02	2	8.40E+03	18	0.17	0.8490

Table 6. Results from two-factor analysis of variance for T₃ percent change

Exposure Day	Source	SS	DF	SSE	DFE	F	P
	Dose	1.17E+02	2	1.68E+03	17	0.59	0.5643
2	Sex	1.68E+02	1	1.68E+03	17	1.70	0.2103
	Dose*Sex	2.20E+02	2	1.68E+03	17	1.11	0.3525
	Dose	8.25E+01	2	1.96E+03	18	0.38	0.6901
3	Sex	2.36E+01	1	1.96E+03	18	0.22	0.6470
	Dose*Sex	3.07E+02	2	1.96E+03	18	1.41	0.2699
	Dose	3.85E+02	2	1.97E+03	17	1.66	0.2196
4	Sex	1.42E+01	1	1.97E+03	17	0.12	0.7308
	Dose*Sex	6.39E+02	2	1.97E+03	17	2.76	0.0918
	Dose	3.83E+02	2	1.87E+03	9	0.92	0.4315
8	Sex	4.37E+00	1	1.87E+03	9	0.02	0.8878
<u> </u>	Dose*Sex	5.07E+02	2	1.87E+03	9	1.22	0.3390
	Dose	2.09E+02	2	2.83E+03	17	0.63	0.5449
14	Sex	1.49E+02	1	2.83E+03	17	0.89	0.3574
	Dose*Sex	3.68E+02	2	2.83E+03	17	1.11	0.3534
	Dose	6.77E+01	2	4.34E+03	18	0.14	0.8701
Post 1	Sex	2.14E+01	1	4.34E+03	18	0.09	0.7691
	Dose*Sex	6.90E+02	2	4.34E+03	18	1.43	0.2651
	Dose	6.32E+01	2	3.09E+03	17	0.17	0.8422
Post 15	Sex	1.82E+02	1	3.09E+03	17	1.00	0.3310
	Dose*Sex	3.04E+02	2	3.09E+03	17	0.84	0.4508

Figure 2 contains the mean percent change from baseline for each combination of dose group and exposure day. Means were averaged within each sex first and then averaged across sex. When comparing across exposure days within a dose group, keep in mind there were instances of missing data. No attempt was made to estimate missing data since all statistical tests were performed within an exposure day.

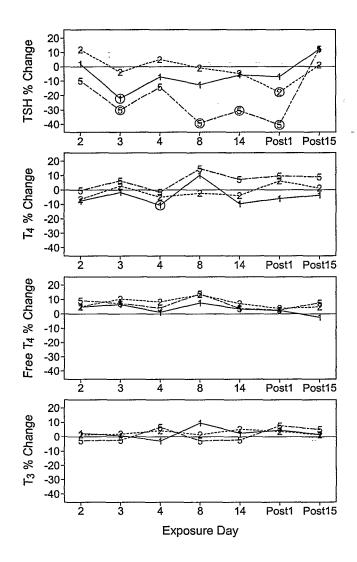


Figure 2. Mean percent change from baseline. Legend: 2 = 0.02 mg/kg-day, 1 = 0.1 mg/kg-day, 5 = 0.5 mg/kg-day. Significant changes from baseline ($p \le 0.05$) are circled.

Figure 3 contains the TSH mean percent change from baseline for each combination of sex, dose group and exposure day. The purpose of Figure 3 is to help interpret the significant Dose*Sex interaction for TSH at post exposure day 1 and 15. Means from exposure day 14 were included to see the change from the last day of exposure into post exposure. Table 7 contains the data from each subject used to determine the means used in Figure 3.

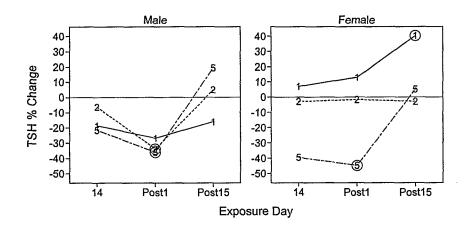


Figure 3. Mean percent change from baseline. Legend: 2 = 0.02 mg/kg-day, 1 = 0.1 mg/kg-day, 5 = 0.5 mg/kg-day. Significant changes from baseline ($p \le 0.05$) are circled.

Table 7. TSH percent change from baseline for each subject

Dose Group			Exposure Day						
(mg/kg-day)	Sex	Subject	14	Post 1	Post 15				
		A	-24	-27	-32				
	M	В	-17	-36	8				
Į		C	38	-28	15				
0.02		D	-23	-45	28				
		E	-13	-20	27				
	F	F	30	20	-45				
		G	-25	-6	- 9				
		H		0	17				
-		I	9	0	9				
1	M	J		-44	-33				
0.1		K	-40	-25	-40				
		L	-25	-38	0				
		M		27	27				
	F	N	6	25	56				
]		0	-10	20	45				
		P	25	-20	33				
		Q	-63	-54	-29				
	M	R	-38	-52	8				
		S	11	-21	63				
0.5		T	4	-17	35				
		U	-65	-61	-25				
	F	V	12	-18	18				
		W	-50	-50	16				
L		X	-55	-51	13				

Figure 4 and Table 8 contain the mean percent change from baseline for each combination of sex, dose group and exposure day. Figures 5 through 8 contain the percent change from baseline for each subject.

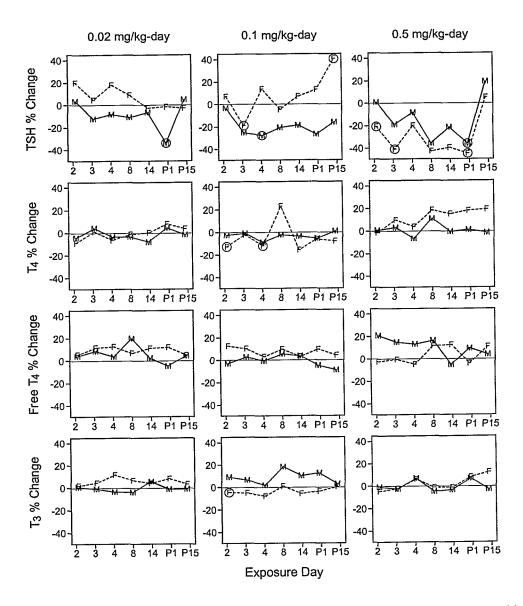


Figure 4. Mean percent change from baseline. Legend: M = Male, F = Female. Significant changes from baseline ($p \le 0.05$) are circled.

Table 8. Mean and standard deviation of subjects for percent change from baseline

Hormone	mg/kg-day	C		Exposure Day							
		Sex	2	3	4	8	14	Post 1	Post 15		
1		M	4±9	-12 ± 27	-8 ± 14	-11 ± 4	-6 ± 30	-34 ± 8	5 ± 26		
	0.02	F	20 ± 33	5 ± 25	18 ± 35	9 ± 23	-3 ± 29	-2 ± 17	-3 ± 32		
L		Avg	12 ± 22	-4 ± 26	5 ± 27	-1 ± 19	-5 ± 30	-18 ± 13	1 ± 29		
		M	-3 ± 20	-25 ± 21	-28 ± 15	-21 ± 10	-19 ± 25	-27 ± 20	-16 ± 24		
TSH	0.1	F	7 ± 14	-19 ± 8	14 ± 50	-5 ± 18	7 ± 18	13 ± 22	40 ± 13		
L		Avg	2 ± 17	-22 ± 16	-7 ± 37	-13 ± 16	-6 ± 22	-7 ± 21	12 ± 19		
		M	1 ± 27	-19 ± 25	-8 ± 22	-36 ± 29	-22 ± 35	-36 ± 20	19 ± 39		
	0.5	F	-21 ± 9	-41 ± 5	-20 ± 21	-43 ± 10	-40 ± 35	-45 ± 19	5 ± 20		
		Avg	-10 ± 20	-30 ± 18	-14 ± 22	-39 ± 24	-31 ± 35	-40 ± 19	12 ± 31		
		M	-4 ± 8	4 ± 22	-3 ± 16	-3 ± 5	-8 ± 6	4 ± 20	-1 ± 8		
0.0	0.02	F	-9 ± 7	1 ± 9	-6 ± 12	-1 ± 8	0 ± 7	8 ± 21	4 ± 15		
Ĺ		Avg	-7 ± 7	3 ± 17	-5 ± 14	-2 ± 7	-4 ± 6	6 ± 20	1 ± 12		
		M	-3 ± 12	-1 ± 20	-9 ± 8	-2 ± 23	-4 ± 16	-6 ± 10	1 ± 6		
T_4	0.1	F	-13 ± 4	-2 ± 11	-12 ± 4	23 ± 24	-16 ± 16	-6 ± 7	-8 ± 11		
		Avg	-8 ± 9	-2 ± 16	-11±6	10 ± 24	-10 ± 16	-6 ± 9	-4 ± 9		
	0.5	M	1 ± 6	3 ± 18	-7 ± 3	11 ± 12	-1 ± 11	1 ± 12	-1 ± 14		
		F	-1 ± 7	10 ± 18	4 ± 11	18 ± 14	15 ± 30	18 ± 21	19 ± 32		
		Avg	0 ± 7	6 ± 18	-2 ± 8	15 ± 13	7 ± 23	10 ± 17	9 ± 24		
	0.02	M	4 ± 16	9 ± 10	4 ± 26	20 ± 0	3 ± 4	-5 ± 13	4 ± 13		
		F	6 ± 10	12 ± 19	13 ± 25	7 ± 14	11 ± 13	12 ± 19	5 ± 16		
L		Avg	5 ± 14	10 ± 15	8 ± 25	13 ± 12	7±9	4 ± 16	5 ± 14		
ſ		M	-3 ± 17	3 ± 18	-1 ± 14	6 ± 18	4 ± 13	-5 ± 11	-9 ± 11		
free T ₄	0.1	F	13 ± 26	10 ± 21	3 ± 14	10 ± 17	3 ± 26	10 ± 32	4 ± 31		
1		Avg	5 ± 22	6 ± 20	1 ± 14	8 ± 18	3 ± 22	2 ± 24	-2 ± 23		
Γ		M	21 ± 21	15 ± 20	13 ± 15	16 ± 21	-5 ± 13	9 ± 16	4 ± 12		
	0.5	F	-3 ± 30	-1 ± 19	-5 ± 10	12 ± 35	13 ± 32	-4 ± 17	11 ± 34		
		Avg	9 ± 26	7 ± 19	4 ± 12	14 ± 27	4 ± 26	3 ± 17	8 ± 26		
		M	1 ± 7	-1 ± 13	-3 ± 13	-4 ± 3	6±10	-1 ± 11	-1 ± 3		
	0.02	F	2 ± 13	5 ± 4	12 ± 16	7 ± 3	4 ± 7	8 ± 13	3 ± 15		
		Avg	1 ± 10	2 ± 9	4 ± 15	1 ± 3	5 ± 8	4 ± 12	1 ± 11		
		М	9 ± 13	7±11	2 ± 8	18 ± 16	10 ± 9	13 ± 9	3 ± 5		
T ₃	0.1	F	-5 ± 2	-5 ± 6	-8 ± 7	1 ± 17	-6 ± 14	-4 ± 6	0 ± 7		
-		Avg	2 ± 10	1 ± 9	-3 ± 8	9 ± 17	2 ± 12	4 ± 8	2 ± 6		
E E		M	-1 ± 9	-3 ± 14	7±4	-5 ± 17	-3 ± 13	7 ± 17	-3 ± 8		
	0.5	F	-5 ± 12	-2 ± 11	6 ± 10	-1 ± 21	-1 ± 19	8 ± 27	13 ± 26		
		Avg	-3 ± 11	-2 ± 13	7±8	-3 ± 18	-2 ± 16	8 ± 23	5 ± 19		

Note: The standard deviation for the Avg row (average of males and females) is the pooled standard deviation across sex.

Although there were missing data in the original raw, there was only one value deleted for being considered an outlier. This was a TSH value for subject M (female, 0.1 mg/kg-day) on exposure day 14 reported as 133% change from baseline.

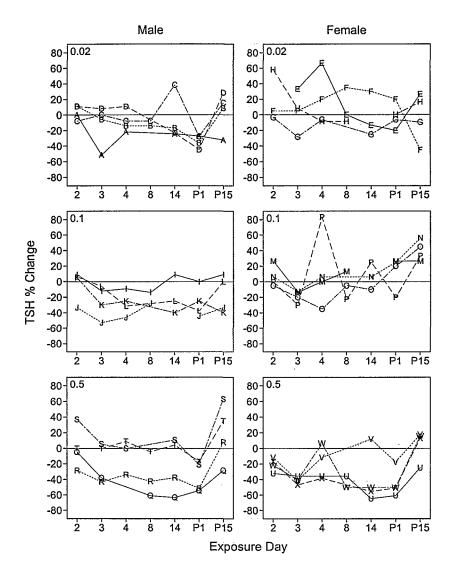


Figure 5. Percent change in TSH from baseline for each subject (4 males and 4 females per dose group). The dose group is identified in the upper left corner of each plot.

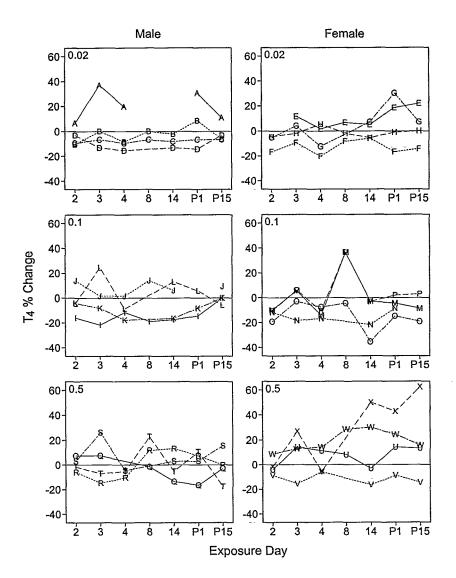


Figure 6. Percent change in T_4 from baseline for each subject (4 males and 4 females per dose group). The dose group is identified in the upper left corner of each plot

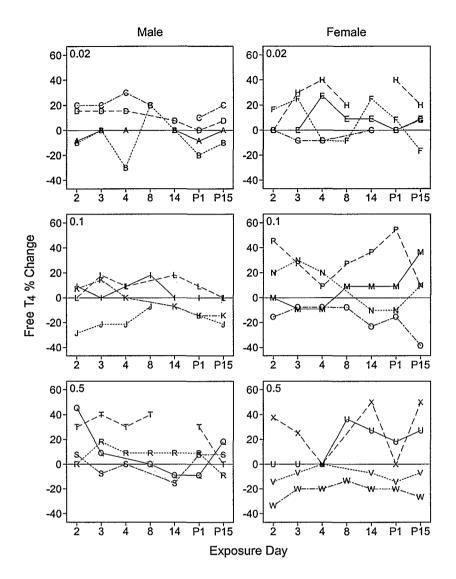


Figure 7. Percent change in free T_4 from baseline for each subject (4 males and 4 females per dose group). The dose group is identified in the upper left corner of each plot.

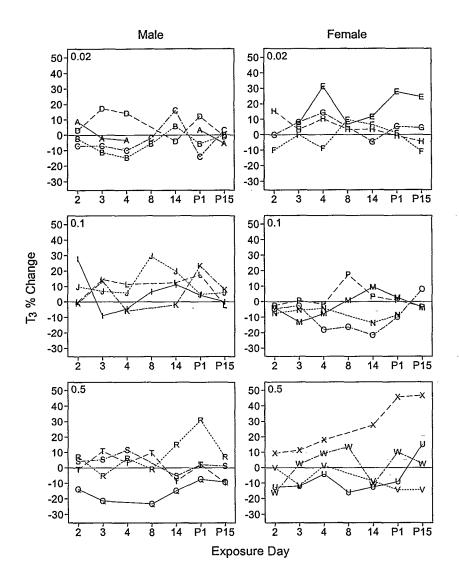


Figure 8. Percent change in T_3 from baseline for each subject (4 males and 4 females per dose group). The dose group is identified in the upper left corner of each plot.

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